

## PATENT COOPERATION TREATY

PCT

From the INTERNATIONAL BUREAU

To:

Commissioner

US Department of Commerce  
 United States Patent and Trademark  
 Office, PCT  
 2011 South Clark Place Room  
 CP2/5C24  
 Arlington, VA 22202  
 ETATS-UNIS D'AMERIQUE

in its capacity as elected Office

## NOTIFICATION OF ELECTION

(PCT Rule 61.2)

Date of mailing (day/month/year)

22 May 2001 (22.05.01)

International application No.

PCT/EP00/07835

Applicant's or agent's file reference

K 2840 Wd

International filing date (day/month/year)

11 August 2000 (11.08.00)

Priority date (day/month/year)

13 August 1999 (13.08.99)

Applicant

NÜESCH, Jürg et al

1. The designated Office is hereby notified of its election made:



in the demand filed with the International Preliminary Examining Authority on:

10 March 2001 (10.03.01)



in a notice effecting later election filed with the International Bureau on:

2. The election ☒ was

was not

made before the expiration of 19 months from the priority date or, where Rule 32 applies, within the time limit under Rule 32.2(b).

The International Bureau of WIPO  
 34, chemin des Colombettes  
 1211 Geneva 20, Switzerland

Authorized officer

Claudio Borton

## (12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization  
International Bureau(43) International Publication Date  
22 February 2001 (22.02.2001)

PCT

(10) International Publication Number  
WO 01/12666 A1(51) International Patent Classification<sup>7</sup>: C07K 14/015[CH/CH]: In den Wegscheiden 1, CH-4132 Muttenz (CH).  
ROMMELAERE, Jean [BE/DE]; Schloss Wolfsbrunn-  
enweg 11, D-69118 Heidelberg (DE).

(21) International Application Number: PCT/EP00/07835

(22) International Filing Date: 11 August 2000 (11.08.2000)

(74) Agent: SCHÜSSLER, Andrea; Huber & Schüssler, Trud-  
eringer Strasse 246, D-81825 München (DE).

(25) Filing Language:

English

(81) Designated States (national): JP, US.

(26) Publication Language:

English

## Published:

(30) Priority Data:

99115161.4

13 August 1999 (13.08.1999) EP

— With international search report.

— Before the expiration of the time limit for amending the  
claims and to be republished in the event of receipt of  
amendments.(71) Applicant (for all designated States except US):  
DEUTSCHES KREBSFORSCHUNGSZENTRUM  
[DE/DE]; Stiftung des öffentlichen Rechts, Im Neuen-  
heimer Feld 280, D-69120 Heidelberg (DE).— With (an) indication(s) in relation to deposited biological  
material furnished under Rule 13bis separately from the  
description.For two-letter codes and other abbreviations, refer to the "Guid-  
ance Notes on Codes and Abbreviations" appearing at the begin-  
ning of each regular issue of the PCT Gazette.

(72) Inventors; and

(75) Inventors/Applicants (for US only): NÜESCH, Jürg

(54) Title: PARVOVIRUS NS1 VARIANTS

(57) Abstract: The present invention relates to a parvovirus NS1 variant having a shifted equilibrium between the DNA replica-  
tion and transcription activities (a) and the cytotoxicity activity (b). Furthermore, this invention relates to DNAs coding for these  
parvovirus NS1 variants and methods of producing them. Additionally, this invention concerns antibodies directed against the par-  
vovirus NS1 variants as well as the use of the DNAs and the parvovirus NS1 variants.

WO 01/12666 A1

**PCT****REQUEST**

For receiving Office use only

International Application No.

International Filing Date

The undersigned requests that the present international application be processed according to the Patent Cooperation Treaty.

Name of receiving Office and "PCT International Application"

Applicant's or agent's file reference (if desired) (12 characters maximum) K 2840 Wd

**Box No. I TITLE OF INVENTION**

Parvovirus NS1 Variants

**Box No. II APPLICANT**

Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country. The country of the address indicated in this Box is the applicant's State (that is, country) of residence if no State of residence is indicated below.)

Deutsches Krebsforschungszentrum  
Stiftung des öffentlichen Rechts  
Im Neuenheimer Feld 280  
69120 Heidelberg

☐ This person is also inventor.

Telephone No.

Facsimile No.

Teleprinter No.

State (that is, country) of nationality:

DE

State (that is, country) of residence:

DE

This person is applicant for the purposes of:

☐ all designated States☒ all designated States except the United States of America☐ the United States of America only☐ the States indicated in the Supplemental Box**Box No. III FURTHER APPLICANT(S) AND/OR (FURTHER) INVENTOR(S)**

Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country. The country of the address indicated in this Box is the applicant's State (that is, country) of residence if no State of residence is indicated below.)

Nüesch, Jürg  
In den Wegscheiden 1  
CH-4132 Muttenz

This person is:

☐ applicant only☒ applicant and inventor☐ inventor only (If this check-box is marked, do not fill in below.)

State (that is, country) of nationality:

CH

State (that is, country) of residence:

CH

This person is applicant for the purposes of:

☐ all designated States☐ all designated States except the United States of America☒ the United States of America only☐ the States indicated in the Supplemental Box☒ Further applicants and/or (further) inventors are indicated on a continuation sheet.**Box No. IV AGENT OR COMMON REPRESENTATIVE; OR ADDRESS FOR CORRESPONDENCE**

The person identified below is hereby/has been appointed to act on behalf of the applicant(s) before the competent International Authorities as:

☒ agent☐ common representative

Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country.)

Dr. Andrea Schüssler

**HUBER & SCHÜSSLER**  
Patentanwälte · Patent Anwälte  
Tuderinger Straße 246 · 81825 München  
Tel. 089/42 72 47 48 · Fax 089/42 72 47 49

Telephone No.

Facsimile No.

Teleprinter No.

☐ Address for correspondence: Mark this check-box where no agent or common representative is/has been appointed and the space above is used instead to indicate a special address to which correspondence should be sent.

Sheet No. .... 2

## Continuation of Box No. III FURTHER APPLICANT(S) AND/OR (FURTHER) INVENTOR(S)

*If none of the following sub-boxes is used, this sheet should not be included in the request.*

Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country. The country of the address indicated in this Box is the applicant's State (that is, country) of residence if no State of residence is indicated below.)

This person is:

Rommelaere, Jean  
Schloß Wolfsbrunnenweg 11  
D-69118 Heidelberg

☐ applicant only☒ applicant and inventor☐ inventor only (If this check-box is marked, do not fill in below.)

State (that is, country) of nationality:

B

State (that is, country) of residence:

DE

This person is applicant for the purposes of:

☐ all designated States☐ all designated States except the United States of America☒ the United States of America only☐ the States indicated in the Supplemental Box

Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country. The country of the address indicated in this Box is the applicant's State (that is, country) of residence if no State of residence is indicated below.)

This person is:

☐ applicant only☐ applicant and inventor☐ inventor only (If this check-box is marked, do not fill in below.)

State (that is, country) of nationality:

State (that is, country) of residence:

This person is applicant for the purposes of:

☐ all designated States☐ all designated States except the United States of America☐ the United States of America only☐ the States indicated in the Supplemental Box

Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country. The country of the address indicated in this Box is the applicant's State (that is, country) of residence if no State of residence is indicated below.)

This person is:

☐ applicant only☐ applicant and inventor☐ inventor only (If this check-box is marked, do not fill in below.)

State (that is, country) of nationality:

State (that is, country) of residence:

This person is applicant for the purposes of:

☐ all designated States☐ all designated States except the United States of America☐ the United States of America only☐ the States indicated in the Supplemental Box

Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country. The country of the address indicated in this Box is the applicant's State (that is, country) of residence if no State of residence is indicated below.)

This person is:

☐ applicant only☐ applicant and inventor☐ inventor only (If this check-box is marked, do not fill in below.)

State (that is, country) of nationality:

State (that is, country) of residence:

This person is applicant for the purposes of:

☐ all designated States☐ all designated States except the United States of America☐ the United States of America only☐ the States indicated in the Supplemental Box☐ Further applicants and/or (further) inventors are indicated on another continuation sheet.

Sheet No. 3

## Box No.V DESIGNATION OF STATES

The following designations are hereby made under Rule 4.9(a) (mark the applicable check-boxes; at least one must be marked):

## Regional Patent

- ☐ AP ARIPO Patent: GH Ghana, GM Gambia, KE Kenya, LS Lesotho, MW Malawi, MZ Mozambique, SD Sudan, SL Sierra Leone, SZ Swaziland, TZ United Republic of Tanzania, UG Uganda, ZW Zimbabwe, and any other State which is a Contracting State of the Harare Protocol and of the PCT
- ☐ EA Eurasian Patent: AM Armenia, AZ Azerbaijan, BY Belarus, KG Kyrgyzstan, KZ Kazakhstan, MD Republic of Moldova, RU Russian Federation, TJ Tajikistan, TM Turkmenistan, and any other State which is a Contracting State of the Eurasian Patent Convention and of the PCT
- ☐ EP European Patent: AT Austria, BE Belgium, CH and LI Switzerland and Liechtenstein, CY Cyprus, DE Germany, DK Denmark, ES Spain, FI Finland, FR France, GB United Kingdom, GR Greece, IE Ireland, IT Italy, LU Luxembourg, MC Monaco, NL Netherlands, PT Portugal, SE Sweden, and any other State which is a Contracting State of the European Patent Convention and of the PCT
- ☐ OA OAPI Patent: BF Burkina Faso, BJ Benin, CF Central African Republic, CG Congo, CI Côte d'Ivoire, CM Cameroon, GA Gabon, GN Guinea, GW Guinea-Bissau, ML Mali, MR Mauritania, NE Niger, SN Senegal, TD Chad, TG Togo, and any other State which is a member State of OAPI and a Contracting State of the PCT (if other kind of protection or treatment desired, specify on dotted line)

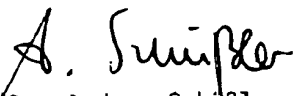
## National Patent (if other kind of protection or treatment desired, specify on dotted line):

- |   |   |
|---|---|
| <input type="checkbox"/> AE United Arab Emirates                  | <input type="checkbox"/> LC Saint Lucia                               |
| <input type="checkbox"/> AG Antigua and Barbuda                   | <input type="checkbox"/> LK Sri Lanka                                 |
| <input type="checkbox"/> AL Albania                               | <input type="checkbox"/> LR Liberia                                   |
| <input type="checkbox"/> AM Armenia                               | <input type="checkbox"/> LS Lesotho                                   |
| <input type="checkbox"/> AT Austria                               | <input type="checkbox"/> LT Lithuania                                 |
| <input type="checkbox"/> AU Australia                             | <input type="checkbox"/> LU Luxembourg                                |
| <input type="checkbox"/> AZ Azerbaijan                            | <input type="checkbox"/> LV Latvia                                    |
| <input type="checkbox"/> BA Bosnia and Herzegovina                | <input type="checkbox"/> MA Morocco                                   |
| <input type="checkbox"/> BB Barbados                              | <input type="checkbox"/> MD Republic of Moldova                       |
| <input type="checkbox"/> BG Bulgaria                              | <input type="checkbox"/> MG Madagascar                                |
| <input type="checkbox"/> BR Brazil                                | <input type="checkbox"/> MK The former Yugoslav Republic of Macedonia |
| <input type="checkbox"/> BY Belarus                               | <input type="checkbox"/> MN Mongolia                                  |
| <input type="checkbox"/> BZ Belize                                | <input type="checkbox"/> MW Malawi                                    |
| <input type="checkbox"/> CA Canada                                | <input type="checkbox"/> MX Mexico                                    |
| <input type="checkbox"/> CH and LI Switzerland and Liechtenstein  | <input type="checkbox"/> MZ Mozambique                                |
| <input type="checkbox"/> CN China                                 | <input type="checkbox"/> NO Norway                                    |
| <input type="checkbox"/> CR Costa Rica                            | <input type="checkbox"/> NZ New Zealand                               |
| <input type="checkbox"/> CU Cuba                                  | <input type="checkbox"/> PL Poland                                    |
| <input type="checkbox"/> CZ Czech Republic                        | <input type="checkbox"/> PT Portugal                                  |
| <input type="checkbox"/> DE Germany                               | <input type="checkbox"/> RO Romania                                   |
| <input type="checkbox"/> DK Denmark                               | <input type="checkbox"/> RU Russian Federation                        |
| <input type="checkbox"/> DM Dominica                              | <input type="checkbox"/> SD Sudan                                     |
| <input type="checkbox"/> DZ Algeria                               | <input type="checkbox"/> SE Sweden                                    |
| <input type="checkbox"/> EE Estonia                               | <input type="checkbox"/> SG Singapore                                 |
| <input type="checkbox"/> ES Spain                                 | <input type="checkbox"/> SI Slovenia                                  |
| <input type="checkbox"/> FI Finland                               | <input type="checkbox"/> SK Slovakia                                  |
| <input type="checkbox"/> GB United Kingdom                        | <input type="checkbox"/> SL Sierra Leone                              |
| <input type="checkbox"/> GD Grenada                               | <input type="checkbox"/> TJ Tajikistan                                |
| <input type="checkbox"/> GE Georgia                               | <input type="checkbox"/> TM Turkmenistan                              |
| <input type="checkbox"/> GH Ghana                                 | <input type="checkbox"/> TR Turkey                                    |
| <input type="checkbox"/> GM Gambia                                | <input type="checkbox"/> TT Trinidad and Tobago                       |
| <input type="checkbox"/> HR Croatia                               | <input type="checkbox"/> TZ United Republic of Tanzania               |
| <input type="checkbox"/> HU Hungary                               | <input type="checkbox"/> UA Ukraine                                   |
| <input type="checkbox"/> ID Indonesia                             | <input type="checkbox"/> UG Uganda                                    |
| <input type="checkbox"/> IL Israel                                | <input checked="" type="checkbox"/> US United States of America       |
| <input type="checkbox"/> IN India                                 | <input type="checkbox"/> UZ Uzbekistan                                |
| <input type="checkbox"/> IS Iceland                               | <input type="checkbox"/> VN Viet Nam                                  |
| <input checked="" type="checkbox"/> JP Japan                      | <input type="checkbox"/> YU Yugoslavia                                |
| <input type="checkbox"/> KE Kenya                                 | <input type="checkbox"/> ZA South Africa                              |
| <input type="checkbox"/> KG Kyrgyzstan                            | <input type="checkbox"/> ZW Zimbabwe                                  |
| <input type="checkbox"/> KP Democratic People's Republic of Korea |   |
| <input type="checkbox"/> KR Republic of Korea                     |   |
| <input type="checkbox"/> KZ Kazakhstan                            |   |

Check-box reserved for designating States which have become party to the PCT after issuance of this sheet:

**Precautionary Designation Statement:** In addition to the designations made above, the applicant also makes under Rule 4.9(b) all other designations which would be permitted under the PCT except any designation(s) indicated in the Supplemental Box as being excluded from the scope of this statement. The applicant declares that those additional designations are subject to confirmation and that any designation which is not confirmed before the expiration of 15 months from the priority date is to be regarded as withdrawn by the applicant at the expiration of that time limit. (Confirmation (including fees) must reach the receiving Office within the 15-month time limit.)

Sheet No. 4

<b>Box No. VI PRIORITY CLAIM</b>		<input type="checkbox"/> Further priority claims are indicated in the Supplemental Box.		
Filing date of earlier application (day/month/year)	Number of earlier application	Where earlier application is:		
		national application: country	regional application:* regional Office	international application: receiving Office
item (1) Att. 13. 1999	99 115 161.4		EPO	
item (2)				
item (3)				
<input checked="" type="checkbox"/> The receiving Office is requested to prepare and transmit to the International Bureau a certified copy of the earlier application(s) (only if the earlier application was filed with the Office which for the purposes of the present international application is the receiving Office) identified above as item(s): <span style="float: right;">1</span>				
<small>* Where the earlier application is an ARIPO application, it is mandatory to indicate in the Supplemental Box at least one country party to the Paris Convention for the Protection of Industrial Property for which that earlier application was filed (Rule 4.10(b)(ii)). See Supplemental Box.</small>				
<b>Box No. VII INTERNATIONAL SEARCHING AUTHORITY</b>				
<b>Choice of International Searching Authority (ISA)</b> (if two or more International Searching Authorities are competent to carry out the international search, indicate the Authority chosen; the two-letter code may be used):		<b>Request to use results of earlier search; reference to that search (if an earlier search has been carried out by or requested from the International Searching Authority):</b>  Date (day/month/year)                      Number                      Country (or regional Office)		
ISA / EP				
<b>Box No. VIII CHECK LIST; LANGUAGE OF FILING</b>				
This international application contains the following number of sheets:  request : 4 description (excluding sequence listing part) : 10 claims : 2 abstract : 1 drawings : 6 sequence listing part of description : 27 Total number of sheets : 50		This international application is accompanied by the item(s) marked below: 1. <input checked="" type="checkbox"/> fee calculation sheet 2. <input type="checkbox"/> separate signed power of attorney 3. <input type="checkbox"/> copy of general power of attorney; reference number, if any: 4. <input type="checkbox"/> statement explaining lack of signature 5. <input checked="" type="checkbox"/> priority document(s) identified in Box No. VI as item(s): 6. <input type="checkbox"/> translation of international application into (language): 7. <input checked="" type="checkbox"/> separate indications concerning deposited microorganism or other biological material 8. <input checked="" type="checkbox"/> nucleotide and/or amino acid sequence listing in computer readable form 9. <input type="checkbox"/> other (specify):		
Figure of the drawings which should accompany the abstract:		Language of filing of the international application:		
<b>Box No. IX SIGNATURE OF APPLICANT OR AGENT</b>				
Next to each signature, indicate the name of the person signing and the capacity in which the person signs (if such capacity is not obvious from reading the request).				
<div style="display: flex; justify-content: space-between; align-items: flex-end;"> <div style="text-align: center;">               Dr. Andrea Schüßler              European Patent Attorney           </div> <div style="text-align: right;">             München, 10. Aug. 2000           </div> </div>				

<b>For receiving Office use only</b>	
1. Date of actual receipt of the purported international application:	2. Drawings:  <input type="checkbox"/> received:  <input type="checkbox"/> not received:
3. Corrected date of actual receipt due to later but timely received papers or drawings completing the purported international application:	
4. Date of timely receipt of the required corrections under PCT Article 11(2):	
5. International Searching Authority (if two or more are competent): ISA /	6. <input type="checkbox"/> Transmittal of search copy delayed until search fee is paid.

<b>For International Bureau use only</b>	
Date of receipt of the record copy by the International Bureau:	
<i>See Notes to the request form</i>	

## PATENT COOPERATION TREATY

## PCT

## INTERNATIONAL SEARCH REPORT

(PCT Article 18 and Rules 43 and 44)

Applicant's or agent's file reference <b>K 2840 Wd</b>	<b>FOR FURTHER ACTION</b> see Notification of Transmittal of International Search Report (Form PCT/ISA/220) as well as, where applicable, item 5 below.	
International application No. <b>PCT/EP 00/ 07835</b>	International filing date (day/month/year) <b>11/08/2000</b>	(Earliest) Priority Date (day/month/year) <b>13/08/1999</b>
Applicant <b>DEUTSCHES KREBSFORSCHUNGSZENTRUM et al.</b>		

This International Search Report has been prepared by this International Searching Authority and is transmitted to the applicant according to Article 18. A copy is being transmitted to the International Bureau.

This International Search Report consists of a total of 4 sheets.



It is also accompanied by a copy of each prior art document cited in this report.

## 1. Basis of the report

- a. With regard to the language, the International search was carried out on the basis of the International application in the language in which it was filed, unless otherwise indicated under this item.



the International search was carried out on the basis of a translation of the international application furnished to this Authority (Rule 23.1(b)).

- b. With regard to any nucleotide and/or amino acid sequence disclosed in the international application, the International search was carried out on the basis of the sequence listing:



contained in the International application in written form.



filed together with the International application in computer readable form.



furnished subsequently to this Authority in written form.



furnished subsequently to this Authority in computer readable form.



the statement that the subsequently furnished written sequence listing does not go beyond the disclosure in the international application as filed has been furnished.



the statement that the information recorded in computer readable form is identical to the written sequence listing has been furnished

2.



Certain claims were found unsearchable (See Box I).

3.



Unity of invention is lacking (see Box II).

## 4. With regard to the title,



the text is approved as submitted by the applicant.



the text has been established by this Authority to read as follows:

## 5. With regard to the abstract,



the text is approved as submitted by the applicant.



the text has been established, according to Rule 38.2(b), by this Authority as it appears in Box III. The applicant may, within one month from the date of mailing of this international search report, submit comments to this Authority.

## 6. The figure of the drawings to be published with the abstract is Figure No.



as suggested by the applicant.



because the applicant failed to suggest a figure.



because this figure better characterizes the invention.



None of the figures.

## INTERNATIONAL SEARCH REPORT

International application No.  
PCT/EP 00/07835**Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)**

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:  
  
Although claims 14 and 15 are directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the NS1 variant.
2. ☐ Claims Nos.:  
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

**Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)**

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.



## INTERNATIONAL SEARCH REPORT

International Application No

PCT/EP 00/07835

## A. CLASSIFICATION OF SUBJECT MATTER

IPC 7 C12N15/35 C07K14/015 C07K16/08 G01N33/569 C12Q1/70  
A61K35/76 A61K48/00

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 7 C12N C07K A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the International search (name of data base and, where practical, search terms used)

EPO-Internal, STRAND, MEDLINE, BIOSIS

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	<p>D LEGENDRE &amp; J ROMMELAERE: "Terminal regions of the NS-1 protein of the parvovirus Minute Virus of Mice are involved in cytotoxicity and promoter trans inhibition" JOURNAL OF VIROLOGY, vol. 66, no. 10, October 1992 (1992-10), pages 5705-5713, XP000867510 AMERICAN SOCIETY FOR MICROBIOLOGY US *mutants pMMBa131 and pULB3201; figure 1 and page 5710 first paragraph*</p> <p style="text-align: center;">---</p> <p style="text-align: center;">-/-</p>	1-5,7-11

☒ Further documents are listed in the continuation of box C.☐ Patent family members are listed in annex

## \* Special categories of cited documents:

- \*A\* document defining the general state of the art which is not considered to be of particular relevance
- \*E\* earlier document but published on or after the International filing date
- \*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- \*O\* document referring to an oral disclosure, use, exhibition or other means
- \*P\* document published prior to the International filing date but later than the priority date claimed

\*T\* later document published after the International filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

\*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

\*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

\*Z\* document member of the same patent family

Date of the actual completion of the International search

19 January 2001

Date of mailing of the international search report

30. 01. 01

Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2  
NL - 2280 HV Rijswijk  
Tel. (+31-70) 340-2040, Tx. 31 651 epo nl,  
Fax (+31-70) 340-3016

Authorized officer

Cupido, M

## INTERNATIONAL SEARCH REPORT

International Application No

PCT/EP 00/07835

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	LI X ET AL: "Mutation of lysine 405 to serine in the parvovirus H-1 NS1 abolishes its function for viral DNA replication, late promoter activation, and cytotoxicity" JOURNAL OF VIROLOGY, vol. 64, no. 10, October 1990 (1990-10), pages 4654-4660, XP000867496 AMERICAN SOCIETY FOR MICROBIOLOGY US page 4656 -page 4657	1,3,7, 9-11
A	J P F NÜESCH ET AL: "Sequence motifs in the replicator protein of parvovirus MVM essential for nicking and covalent attachment to the viral origin: identification of the linking tyrosine" VIROLOGY,US,ACADEMIC PRESS,ORLANDO, vol. 209, no. 1, 10 May 1995 (1995-05-10), pages 122-135, XP002088311 ISSN: 0042-6822 page 127 -page 131	1-13
A	S F COTTMORE ET AL: "The NS1 polypeptide of the murine parvovirus MVM binds to DNA sequences containing the motif (ACCA)2-3" JOURNAL OF VIROLOGY,US,THE AMERICAN SOCIETY FOR MICROBIOLOGY, vol. 69, no. 3, pages 1652-1660, XP002088309 ISSN: 0022-538X page 1658, left-hand column, last paragraph -right-hand column	1-13

The demand must be filed directly with the competent International Preliminary Examining Authority or, if two or more Authorities are competent, with the one chosen by the applicant. The full name or two-letter code of that Authority may be indicated by the applicant on the line below:

IPEA/ \_\_\_\_\_

# PCT

## CHAPTER II

### DEMAND

under Article 31 of the Patent Cooperation Treaty:

The undersigned requests that the international application specified below be the subject of international preliminary examination according to the Patent Cooperation Treaty.

For International Preliminary Examining Authority use only	
Identification of IPEA	Date of receipt of DEMAND
<b>Box No. I IDENTIFICATION OF THE INTERNATIONAL APPLICATION</b>	
Applicant's or agent's file reference K 2840 Wd	
International application No. PCT/EP00/07835	International filing date (day/month/year) Aug. 11, 2000
(Earliest) Priority date (day/month/year) Aug. 13, 1999	
Title of invention Parvovirus NS1 Variants	
<b>Box No. II APPLICANT(S)</b>	
Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country.)	
Deutsches Krebsforschungszentrum Stiftung des öffentlichen Rechts Im Neuenheimer Feld 280 69120 Heidelberg	
Telephone No.:	
Facsimile No.:	
Teleprinter No.:	
State (i.e. country) of nationality: DE	State (i.e. country) of residence: DE
Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country.)	
NÜESCH, Jürg In den Wegscheiden 1 CH-4132 Muttenz	
State (i.e. country) of nationality: CH	State (i.e. country) of residence: CH
Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country.)	
ROMMELAERE, Jean Schlöß Wolfsbrunnenweg 11 D-69118 Heidelberg	
State (i.e. country) of nationality: DE	State (i.e. country) of residence: DE
<input type="checkbox"/> Further applicants are indicated on a continuation sheet.	

Sheet No. 2.

International application No.  
PCT/EP00/07835**Box No. III AGENT OR COMMON REPRESENTATIVE; OR ADDRESS FOR CORRESPONDENCE**The following person is ☒ agent ☐ common representativeand ☒ has been appointed earlier and represents the applicant(s) also for international preliminary examination.☐ is hereby appointed and any earlier appointment of (an) agent(s)/common representative is hereby revoked.☐ is hereby appointed, specifically for the procedure before the International Preliminary Examining Authority, in addition to the agent(s)/common representative appointed earlier.Name and address: (Family name followed by given name; for a legal entity, full official designation.  
The address must include postal code and name of country.)

Dr. Andrea Schübler

**HUBER & SCHÜSSLER**  
Patentanwälte · Patent Attorneys  
Trodeninger Straße 246 · 81825 München  
Tel. 089/42 72 47 48 · Fax 089/42 72 47 49

Telephone No.:

Facsimile No.:

Teleprinter No.:

☐ Mark this check-box where no agent or common representative is/has been appointed and the space above is used instead to indicate a special address to which correspondence should be sent.**Box No. IV STATEMENT CONCERNING AMENDMENTS**

The applicant wishes the International Preliminary Examining Authority\*

(i) ☐ to start the international preliminary examination on the basis of the international application as originally filed.(ii) ☐ to take into account the amendments under Article 34 of☐ the description (amendments attached).☐ the claims (amendments attached).☐ the drawings (amendments attached).(iii) ☐ to take into account any amendments of the claims under Article 19 filed with the International Bureau (a copy is attached).(iv) ☐ to disregard any amendments of the claims made under Article 19 and to consider them as reversed.(v) ☐ to postpone the start of the international preliminary examination until the expiration of 20 months from the priority date unless that Authority receives a copy of any amendments made under Article 19 or a notice from the applicant that he does not wish to make such amendments (Rule 69.1(d)). (This check-box may be marked only where the time limit under Article 19 has not yet expired.)

\* Where no check-box is marked, international preliminary examination will start on the basis of the international application as originally filed or, where a copy of amendments to the claims under Article 19 and/or amendments of the international application under Article 34 are received by the International Preliminary Examining Authority before it has begun to draw up a written opinion or the international preliminary examination report, as so amended.

**Box No. V ELECTION OF STATES**☒ The applicant hereby elects all eligible States (that is, all States which have been designated and which are bound by Chapter II of the PCT) except .....

(If the applicant does not wish to elect certain eligible States, the name(s) or country code(s) of those States must be indicated above.)

Sheet No. 3.

International application No.

PCT/EP00/07835

## Box No. VI CHECK LIST

The demand is accompanied by the following documents for the purposes of international preliminary examination:

- |  |   |        |
|--|---|--------|
| 1. amendments under Article 34                     |   |        |
| description  | : | sheets |
| claims   | : | sheets |
| drawings   | : | sheets |
| 2. letter accompanying amendments under Article 34 | : | sheets |
| 3. copy of amendments under Article 19             | : | sheets |
| 4. copy of statement under Article 19              | : | sheets |
| 5. other (specify):                                | : | sheets |

For International Preliminary

Examining Authority use only

received      not received

<input type="checkbox"/>	<input type="checkbox"/>
<input type="checkbox"/>	<input type="checkbox"/>
<input type="checkbox"/>	<input type="checkbox"/>
<input type="checkbox"/>	<input type="checkbox"/>
<input type="checkbox"/>	<input type="checkbox"/>
<input type="checkbox"/>	<input type="checkbox"/>
<input type="checkbox"/>	<input type="checkbox"/>

The demand is also accompanied by the item(s) marked below:

- |  |  |
|--|--|
| 1. <input type="checkbox"/> separate signed power of attorney      | 4. <input checked="" type="checkbox"/> fee calculation sheet |
| 2. <input type="checkbox"/> copy of general power of attorney      | 5. <input checked="" type="checkbox"/> other (specify):      |
| 3. <input type="checkbox"/> statement explaining lack of signature | cheque   |

## Box No. VII SIGNATURE OF APPLICANT, AGENT OR COMMON REPRESENTATIVE

Next to each signature, indicate the name of the person signing and the capacity in which the person signs (if such capacity is not obvious from reading the demand).

March 9, 2001

*A. Schübler*  
Dr. Andrea Schübler  
European Patent Attorney

For International Preliminary Examining Authority use only

1. Date of actual receipt of DEMAND:

2. Adjusted date of receipt of demand due to CORRECTIONS under Rule 60.1(b):

3. ☐ The date of receipt of the demand is AFTER the expiration of 19 months from the priority date and item 4 or 5, below, does not apply.☐ The applicant has been informed accordingly.4. ☐ The date of receipt of the demand is WITHIN the period of 19 months from the priority date as extended by virtue of Rule 80.5.5. ☐ Although the date of receipt of the demand is after the expiration of 19 months from the priority date, the delay in arrival is EXCUSED pursuant to Rule 82.

For International Bureau use only

Demand received from IPEA on:

## PATENT COOPERATION TREATY

From the  
INTERNATIONAL PRELIMINARY EXAMINING AUTHORITY

PCT

To:

SCHÜSSLER, Andrea  
HUBER & SCHÜSSLER  
Truderinger Strasse 246  
D-81825 München  
ALLEMAGNE

Huber & Schüssler  
Patentanwälte

03. DEZ. 2001

Frist: .....

NOTIFICATION OF TRANSMITTAL OF  
THE INTERNATIONAL PRELIMINARY  
EXAMINATION REPORT  
(PCT Rule 71.1)

Date of mailing  
(day/month/year)

29.11.2001

Applicant's or agent's file reference  
K 2840 Wd

## IMPORTANT NOTIFICATION

International application No.  
PCT/EP00/07835

International filing date (day/month/year)  
11/08/2000

Priority date (day/month/year)  
13/08/1999

Applicant

DEUTSCHES KREBSFORSCHUNGSZENTRUM et al.

1. The applicant is hereby notified that this International Preliminary Examining Authority transmits herewith the international preliminary examination report and its annexes, if any, established on the international application.
2. A copy of the report and its annexes, if any, is being transmitted to the International Bureau for communication to all the elected Offices.
3. Where required by any of the elected Offices, the International Bureau will prepare an English translation of the report (but not of any annexes) and will transmit such translation to those Offices.


## 4. REMINDER

The applicant must enter the national phase before each elected Office by performing certain acts (filing translations and paying national fees) within 30 months from the priority date (or later in some Offices) (Article 39(1)) (see also the reminder sent by the International Bureau with Form PCT/IB/301).

Where a translation of the International application must be furnished to an elected Office, that translation must contain a translation of any annexes to the international preliminary examination report. It is the applicant's responsibility to prepare and furnish such translation directly to each elected Office concerned.

For further details on the applicable time limits and requirements of the elected Offices, see Volume II of the PCT Applicant's Guide.

Name and mailing address of the IPEA/

 European Patent Office - P.B. 5816 Patentlaan 2  
NL-2280 HV Rijswijk - Pays Bas  
Tel. +31 70 340 - 2040 Tx: 31 651 epo nl  
Fax: +31 70 340 - 3016

Authorized officer

Cardenas, C

Tel. +31 70 340-3370





## PATENT COOPERATION TREATY

## PCT

## INTERNATIONAL PRELIMINARY EXAMINATION REPORT

(PCT Article 36 and Rule 70)

Applicant's or agent's file reference K 2840 Wd		<b>FOR FURTHER ACTION</b> See Notification of Transmittal of International Preliminary Examination Report (Form PCT/IPEA/416)	
International application No. PCT/EP00/07835	International filing date (day/month/year) 11/08/2000	Priority date (day/month/year) 13/08/1999	
International Patent Classification (IPC) or national classification and IPC C07K14/015			
Applicant DEUTSCHES KREBSFORSCHUNGSZENTRUM et al.			
<p>1. This international preliminary examination report has been prepared by this International Preliminary Examining Authority and is transmitted to the applicant according to Article 36.</p> <p>2. This REPORT consists of a total of 6 sheets, including this cover sheet.</p> <p><input checked="" type="checkbox"/> This report is also accompanied by ANNEXES, i.e. sheets of the description, claims and/or drawings which have been amended and are the basis for this report and/or sheets containing rectifications made before this Authority (see Rule 70.16 and Section 607 of the Administrative Instructions under the PCT).</p> <p>These annexes consist of a total of 2 sheets.</p>			
<p>3. This report contains indications relating to the following items:</p> <ul style="list-style-type: none"> <li>I <input checked="" type="checkbox"/> Basis of the report</li> <li>II <input type="checkbox"/> Priority</li> <li>III <input checked="" type="checkbox"/> Non-establishment of opinion with regard to novelty, inventive step and industrial applicability</li> <li>IV <input type="checkbox"/> Lack of unity of invention</li> <li>V <input checked="" type="checkbox"/> Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement</li> <li>VI <input type="checkbox"/> Certain documents cited</li> <li>VII <input type="checkbox"/> Certain defects in the international application</li> <li>VIII <input checked="" type="checkbox"/> Certain observations on the international application</li> </ul>			
Date of submission of the demand 10/03/2001		Date of completion of this report 29.11.2001	
Name and mailing address of the international preliminary examining authority:  European Patent Office - P.B. 5818 Patentlaan 2 NL-2280 HV Rijswijk - Pays Bas Tel. +31 70 340 - 2040 Tx: 31 651 epo nl Fax: +31 70 340 - 3016		Authorized officer Cupido, M Telephone No. +31 70 340 3374 	

**INTERNATIONAL PRELIMINARY  
EXAMINATION REPORT**

International application No. PCT/EP00/07835

**I. Basis of the report**

1. With regard to the elements of the international application (*Replacement sheets which have been furnished to the receiving Office in response to an invitation under Article 14 are referred to in this report as "originally filed" and are not annexed to this report since they do not contain amendments (Rules 70.16 and 70.17)*):
- Description, pages:**

1-10 as originally filed

**Claims, No.:**

14,15 as originally filed

1-13 as received on 06/11/2001 with letter of 06/11/2001

**Drawings, sheets:**

1/6-6/6 as originally filed

2. With regard to the **language**, all the elements marked above were available or furnished to this Authority in the language in which the international application was filed, unless otherwise indicated under this item:

These elements were available or furnished to this Authority in the following language: , which is:

- ☐ the language of a translation furnished for the purposes of the international search (under Rule 23.1(b)).
- ☐ the language of publication of the international application (under Rule 48.3(b)).
- ☐ the language of a translation furnished for the purposes of international preliminary examination (under Rule 55.2 and/or 55.3).

3. With regard to any **nucleotide and/or amino acid sequence** disclosed in the international application, the international preliminary examination was carried out on the basis of the sequence listing:

- ☐ contained in the international application in written form.
- ☐ filed together with the international application in computer readable form.
- ☐ furnished subsequently to this Authority in written form.
- ☐ furnished subsequently to this Authority in computer readable form.
- ☐ The statement that the subsequently furnished written sequence listing does not go beyond the disclosure in the international application as filed has been furnished.
- ☐ The statement that the information recorded in computer readable form is identical to the written sequence listing has been furnished.

4. The amendments have resulted in the cancellation of:



**INTERNATIONAL PRELIMINARY  
EXAMINATION REPORT**

International application No. PCT/EP00/07835

- ☐ the description, pages:  
☐ the claims, Nos.:  
☐ the drawings, sheets:

5. ☐ This report has been established as if (some of) the amendments had not been made, since they have been considered to go beyond the disclosure as filed (Rule 70.2(c)):

*(Any replacement sheet containing such amendments must be referred to under item 1 and annexed to this report.)*

6. Additional observations, if necessary:

**III. Non-establishment of opinion with regard to novelty, inventive step and industrial applicability**

1. The questions whether the claimed invention appears to be novel, to involve an inventive step (to be non-obvious), or to be industrially applicable have not been examined in respect of:

- ☐ the entire international application.  
☒ claims Nos. 12,13.

because:

- ☒ the said international application, or the said claims Nos. 12,13 relate to the following subject matter which does not require an international preliminary examination (*specify*):  
**see separate sheet**
- ☐ the description, claims or drawings (*indicate particular elements below*) or said claims Nos. are so unclear that no meaningful opinion could be formed (*specify*):
- ☐ the claims, or said claims Nos. are so inadequately supported by the description that no meaningful opinion could be formed.
- ☐ no international search report has been established for the said claims Nos. .

2. A meaningful international preliminary examination cannot be carried out due to the failure of the nucleotide and/or amino acid sequence listing to comply with the standard provided for in Annex C of the Administrative Instructions:

- ☐ the written form has not been furnished or does not comply with the standard.  
☐ the computer readable form has not been furnished or does not comply with the standard.

**V. Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement**

**INTERNATIONAL PRELIMINARY  
EXAMINATION REPORT**

International application No. PCT/EP00/07835

**1. Statement**

Novelty (N)	Yes:	Claims 4, 10, 11
	No:	Claims 1-3, 5-9
Inventive step (IS)	Yes:	Claims
	No:	Claims 1-11
Industrial applicability (IA)	Yes:	Claims 1-11
	No:	Claims

**2. Citations and explanations  
see separate sheet****VIII. Certain observations on the international application**

The following observations on the clarity of the claims, description, and drawings or on the question whether the claims are fully supported by the description, are made:  
**see separate sheet**

**INTERNATIONAL PRELIMINARY  
EXAMINATION REPORT - SEPARATE SHEET**

International application No. PCT/EP00/07835

**Re Item III****Non-establishment of opinion with regard to novelty, inventive step and industrial applicability**

Claims 12 and 13 relate to subject-matter considered by this Authority to be covered by the provisions of Rule 67.1(iv) PCT. Consequently, no opinion will be formulated with respect to the industrial applicability of the subject-matter of these claims (Article 34(4)(a)(I) PCT).

For the assessment of these claims on the question whether they are industrially applicable, no unified criteria exist in the PCT. The patentability can also be dependent upon the formulation of the claims. The EPO, for example, does not recognize as industrially applicable the subject-matter of claims to the use of a compound in medical treatment, but will allow, however, claims to a known compound for first use in medical treatment and the use of such a compound for the manufacture of a medicament for a new medical treatment.

**Re Item V****Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement****I Document**

The following document has been taken into consideration:

D1: J. Virol. 66, 5705-5713 ( Legendre and Rommelaere, 1992)

**II Novelty**

D1 discloses a number of mutants in the NS1 protein of the parvovirus MVM. According to figure 1, mutants pMMBa131 and pULB3201 are still of cytotoxicity class I, showing that cytotoxicity is maintained although at a reduced level, but still regarded as being highly toxic, see page 5709 last paragraph, whereas P38 transactivation and DNA replication are strongly inhibited. pMMBa131 contains a deletion of amino acids 245-313, pULB3201 produces an NS1 with a deletion of 374 amino acids (from positions

**INTERNATIONAL PRELIMINARY  
EXAMINATION REPORT - SEPARATE SHEET**

International application No. PCT/EP00/07835

167 to 547), and these mutations are therefore located at the positions referred to in claim 5. Consequently, there can be no doubt that D1 describes NS-1 mutants and their nucleic acids with the characteristics of the NS-1 mutants and DNAs claimed in claims 1-3 and 5 -9, and these claims lack novelty in view of Article 33(2) PCT.

**III Inventive step**

1. D1 is regarded as the closest prior art with respect to the question whether the claimed subject-matter involves an inventive step. The problem underlying the present application in view of D1 is the provision of further parvovirus NS1 proteins that can be used as a toxin for treating tumoral diseases.

2. The novel solutions to this problem provided and claimed in claim 4 of the present invention consist of four specific NS1 variants, designated as S283A, T363A, T394A and T463A. According to Table 1, these NS1 variants with the exception of T463A, are still cytotoxic, whereas P38 transactivation and DNA replication are strongly reduced. These variants can be regarded as possible candidates for use as a toxin in antitumour treatments. Hence, the subject-matter regarding NS1 variants, S283A, T363A and T394A is regarded to involve an inventive step as required by Article 33(3) PCT. Subject-matter relating to the non-toxic NS1 variant T463A is regarded not to involve an inventive step.

3. The antibody and kit referred to in claims 10 and 11 are characterised by the protein sequences to which these antibodies are directed. Since these antigen sequences are known from D1, and antibodies directed to known antigens are devoid of an inventive step, claims 10 and 11 also do not involve an inventive step and these claims cannot be accepted in view of Article 33(3) PCT.

**Re Item VIII****Certain observations on the international application**

Claim 6 refers to DNA coding for parvovirus NS1 variants having the following phosphorylation site mutants: S283A, T363A, T394A, or T463A, wherein the DNA comprises the DNA of figure 1. The DNA of figure 1 represents the wild-type NS1. Hence, the claim is contradictory and violates the requirements of Article 6 PCT.

Patented 2001

CLAIMS

EP 0007889

K 2840

## Claims

1. A parvovirus NS1 variant having a shifted equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b), wherein

5 - the activities (a) are reduced and eliminated, respectively, and activity (b) is maintained or increased or

10 - activity (b) is reduced and eliminated, respectively, and the activities (a) are maintained or increased.

2. The parvovirus NS1 variant according to claim 1, wherein one or several phosphorylation sites are mutated.

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3. The parvovirus NS1 variant according to claim 2, wherein the mutations are located at sites 283, 363, 394 and/or 463.

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4. The parvovirus NS1 variant according to claim 2 or 3, namely the NS1 variants S283A, T363A, T394A, and T463A.

5. A DNA, coding for the parvovirus NS1 variant according to any one of claims 1 to 4.

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6. The DNA according to claim 5, wherein the DNA comprises:

(a) the DNA of figure 1,

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(b) a DNA hybridizing with the DNA from (a), said DNA comprising the mutated phosphorylation site of the DNA from (a), or

(c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.

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7. An expression vector, comprising the DNA according to

Patented 12 November 2001

GEMSPAT

06/02-2002

2

claim 5 or 6.

- 
8. A transformant, containing the expression vector according to claim 7.

5

9. A method of producing the parvovirus NS1 variant according to any one of claims 1 to 4, comprising the culturing of the transformant according to claim 8 under suitable conditions.

10

10. An antibody, directed against the parvovirus NS1 variant according to any one of claims 1 to 4.

11. Kit comprising:

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- (a) a parvovirus NS1 variant according to the invention,
- (b) a DNA according to the invention, e.g. an expression vector, particularly a parvovirus,
- (c) an antibody according to the invention, as well as
- (d) conventional auxiliary agents, such as solvents,

20

buffers, carriers, markers and controls,

wherein of components (a) to (d) one or more representatives can be present each.

25

12. Use of the parvovirus NS1 variant according to any one of claims 1 to 4 as a toxin for treating tumoral diseases.

13. Use of the DNA according to claim 7 as a vector for gene therapy.

30

replaced by  
Article 348

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**Claims**

1. A parvovirus NS1 variant having a shifted equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b).
2. The parvovirus NS1 variant according to claim 1, wherein the activities (a) are reduced and eliminated, respectively, and activity (b) is maintained or increased.
3. The parvovirus NS1 variant according to claim 1, wherein activity (b) is reduced and eliminated, respectively, and the activities (a) are maintained or increased.
4. The parvovirus NS1 variant according to any one of claims 1 to 3, wherein one or several phosphorylation sites are mutated.
5. The parvovirus NS1 variant according to claim 4, wherein the mutations are located at sites 283, 363, 394 and/or 463.
6. The parvovirus NS1 variant according to claim 4 or 5, namely the NS1 variants S283A, T363A, T394A, and T463A.
7. A DNA, coding for the parvovirus NS1 variant according to any one of claims 1 to 6.
8. The DNA according to claim 7, wherein the DNA comprises:
  - (a) the DNA of figure 1,
  - (b) a DNA hybridizing with the DNA from (a), said DNA comprising the mutated phosphorylation site of the DNA from (a), or
  - (c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.

9. An expression vector, comprising the DNA according to claim

~~7 or 8.~~

---

10. A transformant, containing the expression vector according to claim 9.

11. A method of producing the parvovirus NS1 variant according to any one of claims 1 to 6, comprising the culturing of the transformant according to claim 10 under suitable conditions.

12. An antibody, directed against the parvovirus NS1 variant according to any one of claims 1 to 6.

13. Kit comprising:

- (a) a parvovirus NS1 variant according to the invention,
- (b) a DNA according to the invention, e.g. an expression vector, particularly a parvovirus,
- (c) an antibody according to the invention, as well as
- (d) conventional auxiliary agents, such as solvents, buffers, carriers, markers and controls,

wherein of components (a) to (d) one or more representatives can be present each.

14. Use of the parvovirus NS1 variant according to any one of claims 1 to 6 as a toxin for treating tumoral diseases.

15. Use of the DNA according to claim 9 as a vector for gene therapy.



---

~~Summary~~

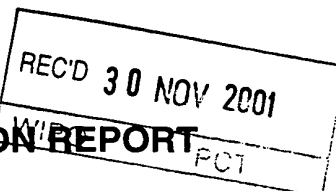
The present invention relates to a parvovirus NS1 variant having a shifted equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b). Furthermore, this invention relates to DNAs coding for these parvovirus NS1 variants and methods of producing them. Additionally, this invention concerns antibodies directed against the parvovirus NS1 variants as well as the use of the DNAs and the parvovirus NS1 variants.

# PATENT COOPERATION TREATY

## PCT

### INTERNATIONAL PRELIMINARY EXAMINATION REPORT

(PCT Article 36 and Rule 70)



2

Applicant's or agent's file reference <b>K 2840 Wd</b>	See Notification of Transmittal of International <b>FOR FURTHER ACTION</b> Preliminary Examination Report (Form PCT/IPEA/416)	
International application No. <b>PCT/EP00/07835</b>	International filing date (day/month/year) <b>11/08/2000</b>	Priority date (day/month/year) <b>13/08/1999</b>
International Patent Classification (IPC) or national classification and IPC <b>C07K14/015</b>		
Applicant <b>DEUTSCHES KREBSFORSCHUNGSZENTRUM et al.</b>		

1. This international preliminary examination report has been prepared by this International Preliminary Examining Authority and is transmitted to the applicant according to Article 36.

2. This REPORT consists of a total of 6 sheets, including this cover sheet.

- ☒ This report is also accompanied by ANNEXES, i.e. sheets of the description, claims and/or drawings which have been amended and are the basis for this report and/or sheets containing rectifications made before this Authority (see Rule 70.16 and Section 607 of the Administrative Instructions under the PCT).

These annexes consist of a total of 2 sheets.

3. This report contains indications relating to the following items:

- I ☒ Basis of the report
- II ☐ Priority
- III ☒ Non-establishment of opinion with regard to novelty, inventive step and industrial applicability
- IV ☐ Lack of unity of invention
- V ☒ Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement
- VI ☐ Certain documents cited
- VII ☐ Certain defects in the international application
- VIII ☒ Certain observations on the international application

Date of submission of the demand  <b>10/03/2001</b>	Date of completion of this report  <b>29.11.2001</b>
Name and mailing address of the international preliminary examining authority:  <b>European Patent Office - P.B. 5818 Patentlaan 2 NL-2280 HV Rijswijk - Pays Bas Tel. +31 70 340 - 2040 Tx: 31 651 epo nl Fax: +31 70 340 - 3016</b>	Authorized officer  <b>Cupido, M</b>  Telephone No. +31 70 340 3374  

# INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No. PCT/EP00/07835

## I. Basis of the report

1. With regard to the **elements** of the international application (*Replacement sheets which have been furnished to the receiving Office in response to an invitation under Article 14 are referred to in this report as "originally filed" and are not annexed to this report since they do not contain amendments (Rules 70.16 and 70.17)*):

### Description, pages:

1-10 as originally filed

### Claims, No.:

14,15 as originally filed

1-13 as received on 06/11/2001 with letter of 06/11/2001

### Drawings, sheets:

1/6-6/6 as originally filed

2. With regard to the **language**, all the elements marked above were available or furnished to this Authority in the language in which the international application was filed, unless otherwise indicated under this item.

These elements were available or furnished to this Authority in the following language: , which is:

- ☐ the language of a translation furnished for the purposes of the international search (under Rule 23.1(b)).
- ☐ the language of publication of the international application (under Rule 48.3(b)).
- ☐ the language of a translation furnished for the purposes of international preliminary examination (under Rule 55.2 and/or 55.3).

3. With regard to any **nucleotide and/or amino acid sequence** disclosed in the international application, the international preliminary examination was carried out on the basis of the sequence listing:

- ☐ contained in the international application in written form.
- ☐ filed together with the international application in computer readable form.
- ☐ furnished subsequently to this Authority in written form.
- ☐ furnished subsequently to this Authority in computer readable form.
- ☐ The statement that the subsequently furnished written sequence listing does not go beyond the disclosure in the international application as filed has been furnished.
- ☐ The statement that the information recorded in computer readable form is identical to the written sequence listing has been furnished.

4. The amendments have resulted in the cancellation of:

**INTERNATIONAL PRELIMINARY  
EXAMINATION REPORT**

International application No. PCT/EP00/07835

- ☐ the description, pages:  
☐ the claims, Nos.:  
☐ the drawings, sheets:

5. ☐ This report has been established as if (some of) the amendments had not been made, since they have been considered to go beyond the disclosure as filed (Rule 70.2(c)):

*(Any replacement sheet containing such amendments must be referred to under item 1 and annexed to this report.)*

6. Additional observations, if necessary:

**III. Non-establishment of opinion with regard to novelty, inventive step and industrial applicability**

1. The questions whether the claimed invention appears to be novel, to involve an inventive step (to be non-obvious), or to be industrially applicable have not been examined in respect of:

- ☐ the entire international application.  
☒ claims Nos. 12,13.

because:

- ☒ the said international application, or the said claims Nos. 12,13 relate to the following subject matter which does not require an international preliminary examination (*specify*):  
**see separate sheet**
- ☐ the description, claims or drawings (*indicate particular elements below*) or said claims Nos. are so unclear that no meaningful opinion could be formed (*specify*):
- ☐ the claims, or said claims Nos. are so inadequately supported by the description that no meaningful opinion could be formed.
- ☐ no international search report has been established for the said claims Nos. .

2. A meaningful international preliminary examination cannot be carried out due to the failure of the nucleotide and/or amino acid sequence listing to comply with the standard provided for in Annex C of the Administrative Instructions:

- ☐ the written form has not been furnished or does not comply with the standard.  
☐ the computer readable form has not been furnished or does not comply with the standard.

**V. Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement**

# INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No. PCT/EP00/07835

## 1. Statement

Novelty (N)	Yes:	Claims	4, 10, 11
	No:	Claims	1-3, 5-9
Inventive step (IS)	Yes:	Claims	
	No:	Claims	1-11
Industrial applicability (IA)	Yes:	Claims	1-11
	No:	Claims	

## 2. Citations and explanations see separate sheet

## VIII. Certain observations on the international application

The following observations on the clarity of the claims, description, and drawings or on the question whether the claims are fully supported by the description, are made:  
see separate sheet

**Re Item III**

**Non-establishment of opinion with regard to novelty, inventive step and industrial applicability**

---

Claims 12 and 13 relate to subject-matter considered by this Authority to be covered by the provisions of Rule 67.1(iv) PCT. Consequently, no opinion will be formulated with respect to the industrial applicability of the subject-matter of these claims (Article 34(4)(a)(I) PCT).

For the assessment of these claims on the question whether they are industrially applicable, no unified criteria exist in the PCT. The patentability can also be dependent upon the formulation of the claims. The EPO, for example, does not recognize as industrially applicable the subject-matter of claims to the use of a compound in medical treatment, but will allow, however, claims to a known compound for first use in medical treatment and the use of such a compound for the manufacture of a medicament for a new medical treatment.

**Re Item V**

**Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement**

**I Document**

The following document has been taken into consideration:

D1: J. Virol. 66, 5705-5713 ( Legendre and Rommelaere, 1992)

**II Novelty**

D1 discloses a number of mutants in the NS1 protein of the parvovirus MVM. According to figure 1, mutants pMMBa131 and pULB3201 are still of cytotoxicity class I, showing that cytotoxicity is maintained although at a reduced level, but still regarded as being highly toxic, see page 5709 last paragraph, whereas P38 transactivation and DNA replication are strongly inhibited. pMMBa131 contains a deletion of amino acids 245-313, pULB3201 produces an NS1 with a deletion of 374 amino acids (from positions

167 to 547), and these mutations are therefore located at the positions referred to in claim 5. Consequently, there can be no doubt that D1 describes NS-1 mutants and their nucleic acids with the characteristics of the NS-1 mutants and DNAs claimed in claims 1-3 and 5 -9, and these claims lack novelty in view of Article 33(2) PCT.

---

### **III Inventive step**

1. D1 is regarded as the closest prior art with respect to the question whether the claimed subject-matter involves an inventive step. The problem underlying the present application in view of D1 is the provision of further parvovirus NS1 proteins that can be used as a toxin for treating tumoral diseases.
2. The novel solutions to this problem provided and claimed in claim 4 of the present invention consist of four specific NS1 variants, designated as S283A, T363A, T394A and T463A. According to Table 1, these NS1 variants with the exception of T463A, are still cytotoxic, whereas P38 transactivation and DNA replication are strongly reduced. These variants can be regarded as possible candidates for use as a toxin in antitumour treatments. Hence, the subject-matter regarding NS1 variants, S283A, T363A and T394A is regarded to involve an inventive step as required by Article 33(3) PCT. Subject-matter relating to the non-toxic NS1 variant T463A is regarded not to involve an inventive step.
3. The antibody and kit referred to in claims 10 and 11 are characterised by the protein sequences to which these antibodies are directed. Since these antigen sequences are known from D1, and antibodies directed to known antigens are devoid of an inventive step, claims 10 and 11 also do not involve an inventive step and these claims cannot be accepted in view of Article 33(3) PCT.

### **Re Item VIII**

#### **Certain observations on the international application**

Claim 6 refers to DNA coding for parvovirus NS1 variants having the following phosphorylation site mutants: S283A, T363A, T394A, or T463A, wherein the DNA comprises the DNA of figure 1. The DNA of figure 1 represents the wild-type NS1. Hence, the claim is contradictory and violates the requirements of Article 6 PCT.

## PCT

## INTERNATIONAL SEARCH REPORT

(PCT Article 18 and Rules 43 and 44)

Applicant's or agent's file reference <b>K 2840 Wd</b>	<b>FOR FURTHER ACTION</b> see Notification of Transmittal of International Search Report (Form PCT/ISA/220) as well as, where applicable, item 5 below.	
International application No. <b>PCT/EP 00/ 07835</b>	International filing date (day/month/year) <b>11/08/2000</b>	(Earliest) Priority Date (day/month/year) <b>13/08/1999</b>
Applicant <b>DEUTSCHES KREBSFORSCHUNGSZENTRUM et al.</b>		

This International Search Report has been prepared by this International Searching Authority and is transmitted to the applicant according to Article 18. A copy is being transmitted to the International Bureau.

This International Search Report consists of a total of 4 sheets.

☒ It is also accompanied by a copy of each prior art document cited in this report.

- Basis of the report**
  - With regard to the **language**, the international search was carried out on the basis of the international application in the language in which it was filed, unless otherwise indicated under this item.
☐ the international search was carried out on the basis of a translation of the international application furnished to this Authority (Rule 23.1(b)).
  - With regard to any **nucleotide and/or amino acid sequence** disclosed in the international application, the international search was carried out on the basis of the sequence listing :
☒ contained in the international application in written form.
☒ filed together with the international application in computer readable form.
☐ furnished subsequently to this Authority in written form.
☐ furnished subsequently to this Authority in computer readable form.
☐ the statement that the subsequently furnished written sequence listing does not go beyond the disclosure in the international application as filed has been furnished.
☐ the statement that the information recorded in computer readable form is identical to the written sequence listing has been furnished
- ☒ **Certain claims were found unsearchable** (See Box I).
- ☐ **Unity of invention is lacking** (see Box II).
- With regard to the **title**,
☒ the text is approved as submitted by the applicant.
☐ the text has been established by this Authority to read as follows:
- With regard to the **abstract**,
☒ the text is approved as submitted by the applicant.
☐ the text has been established, according to Rule 38.2(b), by this Authority as it appears in Box III. The applicant may, within one month from the date of mailing of this international search report, submit comments to this Authority.
- The figure of the **drawings** to be published with the abstract is Figure No.
☐ as suggested by the applicant.
☐ because the applicant failed to suggest a figure.
☐ because this figure better characterizes the invention.
☒ None of the figures.



# INTERNATIONAL SEARCH REPORT

International application No.  
PCT/EP 00/07835

## Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:  
~~Although claims 14 and 15 are directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the NS1 variant.~~
2. ☐ Claims Nos.:  
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## B x II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this International application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

## INTERNATIONAL SEARCH REPORT

International Application No

PCT/EP 00/07835

## A. CLASSIFICATION OF SUBJECT MATTER

IPC 7 C12N15/35 C07K14/015 C07K16/08 G01N33/569 C12Q1/70  
A61K35/76 A61K48/00

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 7 C12N C07K A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

EPO-Internal, STRAND, MEDLINE, BIOSIS

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	D LEGENDRE & J ROMMELAERE: "Terminal regions of the NS-1 protein of the parvovirus Minute Virus of Mice are involved in cytotoxicity and promoter trans inhibition" JOURNAL OF VIROLOGY, vol. 66, no. 10, October 1992 (1992-10), pages 5705-5713, XP000867510 AMERICAN SOCIETY FOR MICROBIOLOGY US *mutants pMMBa131 and pULB3201; figure 1 and page 5710 first paragraph* --- -/--	1-5,7-11



Further documents are listed in the continuation of box C.



Patent family members are listed in annex.

## \* Special categories of cited documents:

- \*A\* document defining the general state of the art which is not considered to be of particular relevance
- \*E\* earlier document but published on or after the international filing date
- \*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- \*O\* document referring to an oral disclosure, use, exhibition or other means
- \*P\* document published prior to the international filing date but later than the priority date claimed

- \*T\* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
- \*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
- \*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.
- \*G\* document member of the same patent family

Date of the actual completion of the international search

19 January 2001

Date of mailing of the international search report

30. 01. 01

Name and mailing address of the ISA

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Authorized officer

Cupido, M

## INTERNATIONAL SEARCH REPORT

International Application No.

PCT/EP 00/07835

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	LI X ET AL: "Mutation of lysine 405 to serine in the parvovirus H-1 NS1 abolishes its function for viral DNA replication, late promoter activation, and cytotoxicity" JOURNAL OF VIROLOGY, vol. 64, no. 10, October 1990 (1990-10), pages 4654-4660, XP000867496 AMERICAN SOCIETY FOR MICROBIOLOGY US page 4656 -page 4657	1,3,7, 9-11
A	J P F NÜESCH ET AL: "Sequence motifs in the replicator protein of parvovirus MVM essential for nicking and covalent attachment to the viral origin: identification of the linking tyrosine" VIROLOGY,US,ACADEMIC PRESS,ORLANDO, vol. 209, no. 1, 10 May 1995 (1995-05-10), pages 122-135, XP002088311 ISSN: 0042-6822 page 127 -page 131	1-13
A	S F COTTMORE ET AL: "The NS1 polypeptide of the murine parvovirus MVM binds to DNA sequences containing the motif (ACCA)2-3" JOURNAL OF VIROLOGY,US,THE AMERICAN SOCIETY FOR MICROBIOLOGY, vol. 69, no. 3, pages 1652-1660, XP002088309 ISSN: 0022-538X page 1658, left-hand column, last paragraph -right-hand column	1-13

### Parvovirus NS1 Variants

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The present invention relates to parvovirus NS1 variants, DNAs coding for them and methods of producing the parvovirus NS1 variants. Furthermore, this invention concerns antibodies directed against the parvovirus NS1 variants as well as the use of the DNAs and the parvovirus NS1 variants.

Parvovirus designates a genus of the virus family Parvoviridae. The parvovirus genus comprises a number of small, icosaedric viruses that can replicate in the absence of a helper virus. Parvovirus contains a single-stranded DNA having a length of about 5.000 bp. At the 3' and 5' ends of the DNA there is one palindromic sequence each. The DNA codes for two capsid proteins, VP1 and VP2, as well as for two regulatory non-structure proteins, NS-1 and NS-2. The latter proteins are phosphorylated and show nuclear or both cytoplasmic and nuclear localization, respectively. NS1 is necessary for viral DNA replication and participates in the regulation of viral gene expression. Particularly, NS1 transactivates the promoter P38 and exhibits DNA-binding, helicase and DNA-nicking activities. Furthermore, NS1 induces cytotoxic and/or cytostatic stress in sensitive host cells.

Parvoviruses are usually well-tolerated by populations of their natural host, in which they persist without apparent pathological signs. This is due to both the protection of foetuses and neonates by maternal immunity, and the striking restriction of parvovirus replication to a narrow range of target proliferating tissues in adult animals. This host tolerance concerns especially rodent parvoviruses, for example the minute virus of mice (MVM) and H-1 virus in their respective natural hosts, namely mice and rats. In addition, humans can be infected with the latter viruses, without any evidence of associated deleterious effects from existing

epidemiological studies and clinical trials. On the other hand, it is known that certain parvoviruses, and especially rodent parvoviruses, are both oncotropic, i.e. accumulate preferentially in neoplastic versus normal tissues, and  
5 oncosuppressive, i.e. have a tumor-suppressive effect towards tumor cells, in various animal models. At least part of the oncosuppressive effect is thought to be due to a direct oncolytic action mediated by NS1. This oncosuppressive effect was also demonstrated against human tumor cells transplanted  
10 in recipient animals.

It is considered to use parvoviruses for therapeutic purposes. On the one hand, it seems to be of interest to use parvoviruses as vectors for therapeutic genes, i.e. for  
15 introducing such genes into the genome of cells. On the other hand, it is considered to use NS1 of parvoviruses as a toxin for treating tumoral diseases. However, initial experiments showed unsatisfactory results.

20 Therefore, it is the object of the present invention to provide a product by which parvoviruses and NS1 thereof, respectively, can be used for the above purposes.

25 According to the invention this is achieved by the subject matters defined in the claims.

The present invention is based on the applicant's findings that it is possible to interfere with the activities of parvovirus NS1 so as to shift the equilibrium existing between  
30 the DNA replication and transcription activities (a) and the cytotoxicity activity (b). In particular, he produced parvovirus NS1 variants in which the DNA replication and transcription activities (a) are reduced and eliminated, respectively, whereas the cytotoxicity activity (b) is  
35 maintained or raised. Moreover, he produced parvovirus NS1 variants in which the cytotoxicity activity (b) is reduced and eliminated, respectively, whereas the DNA replication and transcription activities (a) are maintained or raised.

Examples of such parvovirus NS1 variants are indicated in Table 1 and figure 1. In addition, the applicant recognized that the above parvovirus NS1 variants and expression vectors coding for them, particularly parvoviruses, respectively, are suitable for therapeutic purposes.

According to the invention, the applicant's findings are used to provide a parvovirus NS1 variant in which the equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b) is shifted.

The expression "parvovirus" comprises any parvovirus, particularly a rodent parvovirus, such as minute virus of mice (MVM) and H-1 virus.

The expression "the equilibrium between the DNA replication and transcription activities (a) and the cytotoxicity activity (b) is shifted" refers to the fact that in a parvovirus NS1 variant according to the invention such an equilibrium is shifted as compared to the parvovirus NS1 wild-type. In particular, the equilibrium can be shifted to the effect that the DNA replication and transcription activities (a) are reduced and eliminated, respectively, whereas the cytotoxicity activity (b) is maintained or raised. The cytotoxicity activity (b) can also be reduced and eliminated, respectively, whereas the DNA replication and transcription activities (a) are maintained or raised. Such an equilibrium can be determined by various methods. As regards the determination of the DNA replication activity, reference is made e.g. to methods described in Legendre and Rommelaere, 1992, J. Virol. 66, 5705; Cotmore et al., 1992, Virology 190, 365; Cotmore et al., 1993, J. Virol. 67, 1579, Cotmore and Tattersall, 1994, Embo. J. 13, 4145. As to the determination of the transcription activity reference is made to methods described e.g. in Rhode and Richards, 1987, J. Virol. 61, 2807. Regarding the determination of the cytotoxicity activity reference is made to the below examples.

According to the invention parvovirus NS1 variants are preferred in which the shift of equilibrium is achieved by mutation of one or several phosphorylation sites. Particularly preferred are parvovirus NS1 variants which have a mutation at  
5 ~~one or several of the phosphorylation sites 283, 363, 394 and~~  
463. Even more preferred are the parvovirus NS1 variants S283A, T363A, T394A and T463A, which are indicated in Table 1 and figure 1. In S283A, a serine is exchanged by an alanine at position 283, in T363A a threonine is exchanged by alanine at  
10 position 363, in T394A a threonine is exchanged by alanine at position 394 and in T 463A a threonine is exchanged by alanine at position 463.

A further subject matter of the present invention relates to a  
15 nucleic acid, particularly a DNA, which codes for an above parvovirus NS1 variant. Such a DNA comprises preferably:

- (a) the DNA of fig. 1.1, 1.2, 1.3 and 1.4, respectively
- (b) a DNA hybridizing with the DNA from (a), said DNA  
20 comprising the mutated phosphorylation site of the DNA from (a), or
- (c) a DNA related to the DNA from (a) or (b) via the degenerated genetic code.

25 The DNA of (a) was deposited with DSMZ (Deutsche Sammlung von Mikroorganismen and Zellkulturen) on Aug. 11, 1999, i.e. fig. 1.1 as Escherichia coli pRSV-NS:S283A under DSM 12994, fig. 1.2 as Escherichia coli pRSV-NS:T363A under DSM 12995, fig. 1.3. as Escherichia coli pRSV-NS:T394A under DSM 12996 and  
30 fig. 1.4 as Escherichia coli pRSV-NS:T463A under DSM 12997.

The expression "hybridizing DNA" refers to a DNA which hybridizes with a DNA from (a) under normal conditions, particularly at 20(C below the melting point of the DNA. In  
35 this connection, the expression "hybridizing" refers to conventional hybridization conditions, preferably to hybridization conditions where 5xSSPE, 1 % SDS, 1xDenhardt's solution are used as solution and the hybridization

temperatures are between 35(C and 70(C, preferably 65(C. The hybridization is followed by a wash step first carried out with 2xSSC, 1 % SDS and then with 0.2xSSC at temperatures between 35(C and 70(C, preferably at 65(C. Furthermore,  
5 reference is made to Sambrook et al., Molecular Cloning: A Laboratory Manual, 2nd edition, Cold Spring Harbor Laboratory Press, cold Spring Harbor NY (1989).

10 A DNA according to the invention can be present in a vector and expression vector, respectively. A person skilled in the art is familiar with examples thereof. In the case of an expression vector for E. coli these are e.g. pGEMEX, pUC derivatives, pGEX-2T, pET3b, T7 based expression vectors and pQE-8. For the expression in yeast, e.g. pY100 and Ycpad1 have  
15 to be mentioned while e.g. pKCR, pEFBOS, cDM8, pMSCND, and pCEV4 have to be indicated for the expression in animal cells. The baculovirus expression vector pAcSGHisNT-A is especially suitable for the expression in insect cells.

20 In a preferred embodiment, the vector containing the DNA according to the invention is a virus, e.g. an adenovirus, vaccinia virus, an AAV virus or a parvovirus, such as MVM or H-1, a parvovirus being preferred. The vector may also be a retrovirus, such as MoMULV, MoMuLV, HaMuSV, MuMTV, RSV or  
25 GaLV.

For constructing expression vectors which contain the DNA according to the invention, it is possible to use general methods known in the art. These methods include e.g. in vitro  
30 recombination techniques, synthetic methods and in vivo recombination methods as described in Sambrook et al., supra, for example.

35 Furthermore, the present invention relates to host cells which contain the above described vectors. These host cells include bacteria, yeast, insect and animal cells, preferably mammalian cells. The E. coli strains HB101, DH1, x1776, JM101, JM109, BL21, XL1Blue and SG 13009, the yeast strain



Saccharomyces cerevisiae and the animal cells L, A9, 3T3, FM3A, CHO, COS, Vero, HeLa and the insect cells sf9 are preferred. Methods of transforming these host cells, of phenotypically selecting transformants and of expressing the

5 DNA according to the invention by using the above described vectors are known in the art.

Moreover, the present invention relates to antibodies which specifically recognize an above describe parvovirus NS1  
10 variant, i.e. the region of the parvovirus NS1 variant where the mutation responsible for the shifted equilibrium, particularly a mutated phosphorylation site, is located. The antibodies can be monoclonal, polyclonal or synthetic  
15 antibodies or fragments thereof, e.g. Fab, Fv or scFV fragments. Preferably monoclonal antibodies are concerned. For the production it is favorable to immunize animals - particularly rabbits or chickens for a polyclonal antibody and mice for a monoclonal antibody - with an above parvovirus NS1  
20 variant or with fragments thereof. Further boosters of the animals can be effected with the same parvovirus NS1 variant or with fragments thereof. The polyclonal antibody can then be obtained from the animal serum and egg yolk, respectively. The monoclonal antibody can be obtained according to standard  
25 methods, reference being made particularly to the method by Kohler and Milstein (Nature 256 (1975), 495) and Galfr  (Meth. Enzymol. 73 (1981), 3). In this case, mouse myeloma cells are fused with spleen cells originating from the immunized animals. Antibodies according to the invention can be used in many ways, e.g. for the immunoprecipitation of the above  
30 described parvovirus NS1 variants or for the isolation thereof. The antibodies can be bound in immunoassays in liquid phase or to a solid carrier. In this connection, the antibodies can be labeled in various ways. The person skilled in the art is familiar with suitable markers and labeling  
35 methods. Examples of immunoassays are ELISA and RIA.

The present invention provides parvovirus NS1 variants in which the equilibrium between the DNA replication and

transcription activities (a) and the cytotoxicity activity (b) is shifted. In particular, parvovirus NS1 variants are provided which have a reduced or no cytotoxicity activity, whereas the DNA replication and transcription activities are maintained or increased. ~~Parvovirus NS1 variants are also~~ provided in which the DNA replication and transcription activities are reduced and eliminated, respectively, whereas the cytotoxicity activity is maintained or raised. Thus, the present invention provides products which are suitable for therapeutic purposes. In particular, expression vectors according to the invention, e.g. parvoviruses, can be used for gene-therapeutic measures. Moreover, parvoviruses NS1 variants according to the invention are suitable as toxins, e.g. for treating tumoral diseases.

Therefore, a kit is also provided for the application of the present invention. This kit comprises the following:

- (a) a parvovirus NS1 variant according to the invention,
- (b) a DNA according to the invention, e.g. an expression vector, particularly a parvovirus,
- (c) an antibody according to the invention, as well as
- (d) conventional auxiliary agents, such as solvents, buffers, carriers, markers and controls.

Of component (a) to (d) one or more representatives can be present each.

Brief description of the drawings

Fig. 1 shows the DNA and amino acid sequences of parvovirus NS1 variants according to the invention (fig. 1.1, 1.2, 1.3 and 1.4) as compared to a parvovirus NS1 wild-type. In this connection, the mutated sites in the parvovirus NS1 variants according to the invention are labeled each.

The present invention is explained by the examples.

**Example 1: Preparation and purification of NS1 variants according to the invention**

5  
The DNA of the NS1 variant S283A according to the invention was provided as an EcoRV to BstEII fragment obtained by chimeric PCR using two mutagenic primers. This fragment was then inserted into the corresponding cleaved expression vector  
10 pTHisNS1 (Nuesch et al., Virology 209, (1995), 122) to obtain pTHis NS1:S283A. Such a vector codes for a fusion protein comprising 6 histidine residues (N terminus partner) and S283A of Fig. 1 (C terminus partner). For expression and  
15 purification of S283A the NS1 gene under control of the bacteriophage T7 promoter was transferred into vaccinia virus and expressed in eucaryotic cells by double infection together with vTF7-3 (a vaccinia virus expressing the bacteriophage T7 DNA polymerase). 18 hrs post infection cells were harvested and nuclear extracts prepared. The histidine tagged S283A was  
20 then purified by affinity chromatography on Ni-NTA agarose and analyzed by 10 % SDS-PAGE (Nuesch et al., supra).

It showed that a parvovirus NS1 variant according to the invention can be prepared in highly pure form.

25 The NS1 variants T363A, T394A, and T463A were also produced and purified in the same way.

**Example 2: Preparation and detection of an antibody according to the invention**

30 Tubes were coated with purified NS1 variants prepared as in example 1 and monoclonal antibodies (e.g. scFv) specifically binding to S283A were isolated from human synthetic VH+VL scFV phage library (Griffith et al., EMBO J., 13, (1994), 3245)  
35 according to standard panning protocols after >5 isolation and amplification procedures. The variable region of the isolated scFv harbored in the phagemid were sequenced to identify NS1

variant interacting partner proteins harboring such binding motifs from comparison with known genes in the gene bank.

It showed that monoclonal antibodies according to the invention can be isolated.

In addition, the NS1 variants were used for immunization of animals in order to obtain poly- or monoclonal antibodies.

10

**Example 3: Characterization of the parvovirus NS1 variants S283A, T363A, T394A and T463A according to the invention**

15 The characterization of the parvovirus NS1 variants comprised the determination of the DNA replication, transcription, cytotoxicity, DNA binding, nicking and helicase activities. Known methods were used for this purpose (cf. description, supra). As regards the determination of the helicase activity  
20 reference is made to Stahl et al. 1986, EMBO J. 5, 1999. As to the determination of the nicking activity reference is made to Christensen et al., 1997, J. Virol. 71, 1405 and Nuesch et al., 1995, supra. Regarding the determination of the DNA  
25 binding reference is made to Cotmore et al. 1995, J. Virol. 69, 1652. As far as the determination of the cytotoxicity activity is concerned, the following steps were carried out: NS1 variants were transferred into an expression vector containing the NS1 gene under the control of the parvovirus MVMP4 promoter (genuine promoter driving the non-structural  
30 genes of MVM), and the green fluorescent protein (EGFP) under control of an additional promoter. These constructs were then transfected into A9 cells using lipofectamine (GibcoBRL) according to the manufacturer's instruction and the impact of the NS1 variant on the viability of the cells tested in time  
35 course experiments. Transfected cells were identified by fluorescence of the EGFP. Toxic effects were determined in comparison to wild type NS1 or a vector containing no NS1 gene as a function of time as well as a measure of cytopathic

changes on the cell morphology.

The data indicated in Table 1 were obtained:

5

Table 1

	S283A	T363A	T394A	T463A	wt
P38-TA	+	-	-	++++	++++
ACCA	+	++++	++	++	++
Nick-1	+	-	-	+++	+++
Nick-2	+++	-	-	++++	++++
Nick-3	++	-	-		++++
Heli	++	-	(+)	++++	++++
Rep	+	-	-	+	++++
Cyto	++++++	++	+++	(+)	+++

10

15

**Example 4: NS1 variants' expression after transduction using recombinant viral vectors**

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NS1 expression cassettes containing the NS1 variants according to the invention under control of the parvoviral P4 promoter and a 3'untranslated region from parvovirus MVM to ensure stability and translation of the gene product, were transferred either in a parvovirus genome background as exemplified in example 3, or a heterologous viral genome background, such as vaccinia virus (example 1) or adenovirus. Promoter and terminator regions were exchanged according to the requirements. The nucleic acids containing the NS1 variants were then packaged either in vivo (after transient transfection into eucaryotic cells) or in vitro and the packaged transducing particles were isolated. These transducing units containing NS1 variants were used either for studies concerning gene regulation in tissue culture or animals, but also as therapeutic agents either alone or in combination with other agents (such as cytokines) in gene and cancer therapy approaches.

**Claims**

1. A parvovirus NS1 variant having a shifted equilibrium  
5 ~~between the DNA replication and transcription activities~~  
(a) and the cytotoxicity activity (b).
2. The parvovirus NS1 variant according to claim 1, wherein  
the activities (a) are reduced and eliminated,  
10 respectively, and activity (b) is maintained or  
increased.
3. The parvovirus NS1 variant according to claim 1, wherein  
activity (b) is reduced and eliminated, respectively, and  
15 the activities (a) are maintained or increased.
4. The parvovirus NS1 variant according to any one of claims  
1 to 3, wherein one or several phosphorylation sites are  
20 mutated.
5. The parvovirus NS1 variant according to claim 4, wherein  
the mutations are located at sites 283, 363, 394 and/or  
463.
6. The parvovirus NS1 variant according to claim 4 or 5,  
25 namely the NS1 variants S283A, T363A, T394A, and T463A.
7. A DNA, coding for the parvovirus NS1 variant according to  
any one of claims 1 to 6.  
30
8. The DNA according to claim 7, wherein the DNA comprises:
  - (a) the DNA of figure 1,
  - (b) a DNA hybridizing with the DNA from (a), said  
35 DNA comprising the mutated phosphorylation site of  
the DNA from (a), or
  - (c) a DNA related to the DNA from (a) or (b) via  
the degenerated genetic code.

9. An expression vector, comprising the DNA according to claim 7 or 8.

10. A transformant, containing the expression vector according to claim 9.

---

11. A method of producing the parvovirus NS1 variant according to any one of claims 1 to 6, comprising the culturing of the transformant according to claim 10 under suitable conditions.

12. An antibody, directed against the parvovirus NS1 variant according to any one of claims 1 to 6.

13. Kit comprising:

- (a) a parvovirus NS1 variant according to the invention,
- (b) a DNA according to the invention, e.g. an expression vector, particularly a parvovirus,
- (c) an antibody according to the invention, as well as
- (d) conventional auxiliary agents, such as solvents, buffers, carriers, markers and controls,

wherein of components (a) to (d) one or more representatives can be present each.

14. Use of the parvovirus NS1 variant according to any one of claims 1 to 6 as a toxin for treating tumoral diseases.

15. Use of the DNA according to claim 9 as a vector for gene therapy.

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Fig. 1

## Wild-type NS1

```
ATGGCTGGAAATGCTTACTCTGATGAAGTTTTGGGAGCAACCAACTGGTTAAAGGAAAAA
261 -----+-----+-----+-----+-----+-----+-----+ 320
TACCGACCTTTACGAATGAGACTACTTCAAAACCCCTCGTTGGTTGACCAATTCCTTTTT
M A G N A Y S D E V L G A T N W L K E K -
AGTAACCAGGAAGTGTTCATTTGTTTTTAAAAATGAAATGTTCAACTGAATGGAAAA
321 -----+-----+-----+-----+-----+-----+-----+ 380
TCATTGGTCCTTCACAAGAGTAAACAAAAATTTTACTTTTACAAGTTGACTTACCTTT
S N Q E V F S F V F K N E N V Q L N G K -
GATATCGGATGGAATAGTTACAAAAAGAGCTGCAGGAGGACGAGCTGAAATCTTTACAA
381 -----+-----+-----+-----+-----+-----+-----+ 440
CTATAGCCTACCTTATCAATGTTTTTCTCGACGTCCTCCTGCTCGACTTTAGAAATGTT
D I G W N S Y K K E L Q E D E L K S L Q -
CGAGGAGCGGAAACTACTTGGGACCAAGCGAGGACATGGAATGGGAACCACAGTGGAT
441 -----+-----+-----+-----+-----+-----+-----+ 500
GCTCCTCGCCTTTGATGAACCCTGGTTTCGCTCCTGTACCTTACCCTTTGGTGTACCTA
R G A E T T W D Q S E D M E W E T T V D -
GAAATGACCAAAAAGCAAGTATTCATTTTTGATTCTTTGGTTAAAAAATGTTTATTTGAA
501 -----+-----+-----+-----+-----+-----+-----+ 560
CTTTACTGGTTTTTCGTTTCATAAGTAAAACTAAGAAACCAATTTTTTACAAATAAACTT
E M T K K Q V F I F D S L V K K C L F E -
GTGCTTAACACAAAGAATATATTTCTGGTGATGTTAATTGGTTTGTGCAACATGAATGG
561 -----+-----+-----+-----+-----+-----+-----+ 620
CACGAATTGTGTTTCTTATATAAAGGACCACTACAATTAACCAAACACGTTGTACTTACC
V L N T K N I F P G D V N W F V Q H E W -
GGAAAAGACCAAGGCTGGCACTGCCATGTACTAATTGGAGGAAAGGACTTTAGTCAAGCT
621 -----+-----+-----+-----+-----+-----+-----+ 680
CCTTTTCTGGTTCCGACCGTGACGGTACATGATTAACCTCCTTTCCTGAAATCAGTTCGA
G K D Q G W H C H V L I G G K D F S Q A -
CAAGGGAAATGGTGGAGAAGGCAACTAAATGTTTACTGGAGCAGATGGTTGGTAACAGCC
681 -----+-----+-----+-----+-----+-----+-----+ 740
GTTCCCTTTACCACCTCTTCCGTTGATTTACAAATGACCTCGTCTACCAACCATTGTCCG
Q G K W W R R Q L N V Y W S R W L V T A -
TGTAATGTGCAACTAACACCAGCTGAAAGAATTAACTAAGAGAAATAGCAGAAGACAAT
741 -----+-----+-----+-----+-----+-----+-----+ 800
```



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Fig. 1 (Fortsetzung I)

ACATTACACGTTGATTGTGGTCGACTTTCTTAATTTGATTCTCTTTATCGTCTTCTGTTA  
C N V Q L T P A E R I K L R E I A E D N -  
GAGTGGGTTACTCTACTTACTTATAAGCATAAGCAAACCAAAAAGACTATACCAAGTGT  
801  
CTCACCCAATGAGATGAATGAATATTTCGTATTTCGTTTGGTTTTTTCTGATATGGTTCACA  
860  
E W V T L L T Y K H K Q T K K D Y T K C -  
GTTCTTTTTGGAAACATGATTGCTTACTATTTTTTAACTAAAAAGAAAATAAGCACTAGT  
861  
-----+-----+-----+-----+-----+-----+ 920  
CAAGAAAAACCTTTGTACTAACGAATGATAAAAAATTGATTTTTCTTTTATTTCGTGATCA  
V L F G N M I A Y Y F L T K K K I S T S -  
CCACCAAGAGACGGAGGCTATTTTCTTAGCAGTGACTCTGGCTGGAAAACTAACTTTTTA  
921  
-----+-----+-----+-----+-----+-----+ 980  
GGTGGTTCTCTGCCTCCGATAAAAGAATCGTCACTGAGACCGACCTTTTGATTGAAAAAT  
P P R D G G Y F L S S D S G W K T N F L -  
AAAGAAGGCGAGCGCCATCTAGTGAGCAAACCTATACACTGATGACATGCGGCCAGAAACG  
981  
-----+-----+-----+-----+-----+-----+ 1040  
TTTCTTCCGCTCGCGGTAGATCACTCGTTTGATATGTGACTACTGTACGCCGGTCTTTGC  
K E G E R H L V S K L Y T D D M R P E T -  
GTTGAAACCACAGTAACCACTGCGCAGGAACTAAGCGCGGCAGAATTCAAACCTAAAAAA  
1041  
-----+-----+-----+-----+-----+-----+ 1100  
CAACTTTGGTGTCTATTGGTGACGCGTCTTTGATTGCGCGCGTCTTAAGTTTGATTTTTT  
V E T T V T T A Q E T K R G R I Q T K K -  
GAAGTTTCTATTAAAACTACACTTAAAGAGCTGGTGATAAAAGAGTAACCTCACCAGAG  
1101  
-----+-----+-----+-----+-----+-----+ 1160  
CTTCAAAGATAATTTTGATGTGAATTTCTCGACCACGTATTTTCTCATTGGAGTGGTCTC  
E V S I K T T L K E L V H K R V T S P E -  
GACTGGATGATGATGCAGCCAGACAGTTACATTGAAATGATGGCTCAACCAGGTGGAGAA  
1161  
-----+-----+-----+-----+-----+-----+ 1220  
CTGACCTACTACTACGTGCGGTCTGTCAATGTAACTTTACTACCGAGTTGGTCCACCTCTT  
D W M M M Q P D S Y I E M M A Q P G G E -  
AACCTGCTGAAAAATACGCTAGAGATTTGTACACTAACTCTAGCCAGAACCAAAACAGCA  
1221  
-----+-----+-----+-----+-----+-----+ 1280  
TTGGACGACTTTTTATGCGATCTCTAAACATGTGATTGAGATCGGTCTTGGTTTTGTCGT  
N L L K N T L E I C T L T L A R T K T A -  
TTTGACTTAATTTTAGAAAAAGCTGAAACCAGCAAACCTAAACCACTTTTCACTGCCTGAC  
1281  
-----+-----+-----+-----+-----+-----+ 1340  
AAACTGAATTAAATCTTTTTTCGACTTTGGTTCGTTTGATTGGTTGAAAAGTGACGGACTG  
F D L I L E K A E T S K L T N F S L P D -  
ACAAGAACCTGCAGAAATTTTGTCTTTTCATGGCTGGAACTATGTTAAAGTTTGCCATGCT  
1341  
-----+-----+-----+-----+-----+-----+ 1400  
TGTTCTTGGACGTCTTAAAAACGAAAAAGTACCGACCTTGATACAATTTCAAACGGTACGA

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Fig. 1 (Fortsetzung II)

1401 ATTTGCTGTGTTTTAAACAGACAAGGAGGCAAAAGAAATACTGTTTTATTTTCATGGACCA  
-----+-----+-----+-----+-----+ 1460  
TAAACGACACAAAATTTGTCTGTTCCTCCGTTTTCTTTATGACAAAATAAAGTACCTGGT  
I C C V L N R Q G G K R N T V L F H C P

---

1461 GCCAGCACAGGCAAATCTATTATTGCACAAGCCATAGCACAGCAGTTGGCAATGTTGGT  
-----+-----+-----+-----+-----+ 1520  
CGGTCGTGTCCGTTTAGATAATAACGTGTTCCGGTATCGTGTTTCGTCAACCGTTACAACCA  
A S T G K S I I A Q A I A Q A V G N V G -  
TGCTATAATGCAGCCAATGTAACTTTCCATTTAATGACTGTACCAACAAGAACTTGATT  
1521 -----+-----+-----+-----+-----+ 1580  
ACGATATTACGTCCGTTACATTTGAAAGGTAAATTACTGACATGGTTGTTCTTGAACATA  
C Y N A A N V N F P F N D C T N K N L I -  
TGGGTAGAAGAAGCTGGTAACTTTGGACAGCAAGTAAACCGTTTAAAGCCATTGCTCT  
1581 -----+-----+-----+-----+-----+ 1640  
ACCCATCTTCTTCGACCATTGAAACCTGTCGTTCAATTGGTCAAATTCGGTAAACGAGA  
W V E E A G N F G Q Q V N Q F K A I C S -  
GGTCAAACCTATTTCGATTGATCAAAAAGGAAAAGGCAGCAAACAGATTGAACCAACACCA  
1641 -----+-----+-----+-----+-----+ 1700  
CCAGTTTGATAAGCGTAACTAGTTTTTCCCTTTTCCGTCGTTTGTCTAAGTTGGTTGTGGT  
G Q T I R I D Q K G K G S K Q I E P T P -  
GTCATCATGACCACAAATGAGAACATTACAGTGGTCAGAATAGGCTGCGAAGAAAGACCA  
1701 -----+-----+-----+-----+-----+ 1760  
CAGTAGTACTGGTGTTTACTCTTGTAATGTCACCAGTCTTATCCGACGTTCTTTCTGGT  
V I M T T N E N I T V V R I G C E E R P -  
GAACACACTCAACCAATCAGAGACAGAATGCTTAACATTATCTAACACATACCTTGCCT  
1761 -----+-----+-----+-----+-----+ 1820  
CTTGTGTGAGTTGGTTAGTCTCTGTCTTACGAATTGTAAGTAGATTGTGTATGGAACGGA  
E H T Q P I R D R M L N I H L T H T L P -  
GGTGACTTTGGTTTGGTTGACAAAATGAATGGCCCATGATTTGTGCTTGGTTGGTAAAG  
1821 -----+-----+-----+-----+-----+ 1880  
CCACTGAAACCAAACCAACTGTTTTTACTTACCGGGTACTAAACACGAACCAACCATTC  
G D F G L V D K N E W P M I C A W L V K -  
AATGGTTACCAATCTACCATGGCAAGCTACTGTGCTAAATGGGGCAAAGTTCCTGATTGG  
1881 -----+-----+-----+-----+-----+ 1940  
TTACCAATGGTTAGATGGTACCGTTTCGATGACACGATTTACCCCGTTTCAAGGACTAACC  
N G Y Q S T M A S Y C A K W G K V P D W -  
TCAGAAAACCTGGGCGGAGCCAAAGGTGCCAACTCCTATAAATTTACTAGGTTCCGGCACGC  
1941 -----+-----+-----+-----+-----+ 2000  
AGTCTTTTGACCCGCTCGGTTTCCACGGTTGAGGATATTTAAATGATCCAAGCCGTCGC  
S E N W A E P K V P T P I N L L G S A R -  
TCACCATTACGACACCGAAAAGTACGCCTCTCAGCCAGAACTATGCACTAACTCCACTT  
2001 -----+-----+-----+-----+-----+ 2060

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Fig. 1 (Fortsetzung III)

S P F T T P K S T P L S Q N Y A L T P L -  
GCATCGGATCTCGAGGACCTGGCTTTAGAGCCTTGGAGCACACCAAATACTCCTGTTGCG  
2061 -----+-----+-----+-----+-----+ 2120  
CGTAGCCTAGAGCTCCTGGACCGAAATCTCGGAACCTCGTGTGGTTTATGAGGACAACGC

---

A S D L E D L A L E P W S T P N T P V A -  
GGCACTGCAGAAACCCAGAACACTGGGGAAGCTGGTTCCAAAGCCTGCCAAGATGGTCAA  
2121 -----+-----+-----+-----+-----+ 2180  
CCGTGACGTCTTTGGGTCTTGTGACCCCTTCGACCAAGGTTTCGGACGGTTCTACCAGTT

G T A E T Q N T G E A G S K A C Q D G Q -  
CTGAGCCCAACTTGGTTCAGAGATCGAGGAGGATTTGAGAGCGTGCTTCGGTGCGGAACCG  
2181 -----+-----+-----+-----+-----+ 2240  
GACTCGGGTTGAACCAGTCTCTAGCTCCTCCTAACTCTCGCACGAAGCCACGCCTTGGC

L S P T W S E I E E D L R A C F G A E P -  
TTGAAGAAAGACTTCAGCGAGCCGCTGAACTTGGACTAA  
2241 -----+-----+-----+-----+ 2279  
AACTTCTTTCTGAAGTCGCTCGGCGACTTGAACCTGATT

L K K D F S E P L N L D \* -

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Fig. 1.1

1100 - 261 Wildtype-NS1-Sequence

GAAGTTCTCTATTAAACTACACTTAAAGAGCTGGTGCATAAAAGAGTAACCTCACCAGAG  
1101 -----+-----+-----+-----+-----+-----+-----+ 1160  
CTTCAAGATAATTTTGATGTGAATTTCTCGACCACGTATTTTCTCATTGGAGTGGTCTC  
E V **S** I K T T L K E L V H K R V T S P E -  
→ A S283A

1161 - 2279 Wildtype-NS1-Sequence

Fig. 1.2

1340 - 261 Wildtype-NS1-Sequence

ACAAGAACCTGCAGAATTTTGGCTTTTCATGGCTGGAACTATGTTAAAGTTTGCCATGCT  
1341 -----+-----+-----+-----+-----+-----+ 1400  
TGTTCTTGGACGTCTTAAAAACGAAAAGTACCGACCTTGATACAATTTCAAACGGTACGA  
T R **T** C R I F A F H G W N Y V K V C H A -  
→ A T363A

1401 - 2279 Wildtype-NS1-Sequence

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Fig. 1.3

1400 - 261 Wildtype-NS1-Sequence

```
      → 6
      ┌
      │
1401  ATTTGCTGTGTTTTAAACAGACAAGGAGGCAAAAGAAATACTGTTTTATTTCATGGACCA 1460
      -----+-----+-----+-----+-----+-----+
      TAAACGACACAAAATTTGTCTGTTCCGTTTTCTTTATGACAAAATAAAGTACCTGGT
      I  C  C  V  L  N  R  Q  G  G  K  R  N  T  V  L  F  H  G  P  -
      └
      → A T 394 A
```

1461 - 2279 Wildtype-NS1-Sequence

Fig. 1.4

1640 - 261 Wildtype-NS1-Sequence

```
      → 6
      ┌
      │
1641  GGTCAACTATTTCGCATTGATCAAAAAGGAAAAGGCAGCAAACAGATTGAACCAACACCA 1700
      -----+-----+-----+-----+-----+-----+
      CCAGTTTGATAAGCGTAAGTAGTTTTTCCTTTCCGTCGTTTGTCTAACTTGGTTGTGGT
      G  Q  T  I  R  I  D  Q  K  G  K  G  S  K  Q  I  E  P  T  P  -
      └
      → A T 463 A
```

1701 - 2279 Wildtype-NS1-Sequence

## SEQUENCE LISTING

&lt;110&gt; Deutsches Krebsforschungszentrum

&lt;120&gt; Parvovirus NS1 Variants

&lt;130&gt; K 2840

&lt;140&gt; to be assigned

&lt;150&gt; EP 99 115 161.4

&lt;151&gt; 1999-08-13

&lt;160&gt; 18

&lt;170&gt; PatentIn Ver. 2.1

&lt;210&gt; 1

&lt;211&gt; 2019

&lt;212&gt; DNA

&lt;213&gt; Wildtype Parvovirus NS1

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (1)..(2016)

&lt;400&gt; 1

atg gct gga aat gct tac tct gat gaa gtt ttg gga gca acc aac tgg	48
Met Ala Gly Asn Ala Tyr Ser Asp Glu Val Leu Gly Ala Thr Asn Trp	
1 5 10 15	
tta aag gaa aaa agt aac cag gaa gtg ttc tca ttt gtt ttt aaa aat	96
Leu Lys Glu Lys Ser Asn Gln Glu Val Phe Ser Phe Val Phe Lys Asn	
20 25 30	
gaa aat gtt caa ctg aat gga aaa gat atc gga tgg aat agt tac aaa	144
Glu Asn Val Gln Leu Asn Gly Lys Asp Ile Gly Trp Asn Ser Tyr Lys	
35 40 45	
aaa gag ctg cag gag gac gag ctg aaa tct tta caa cga gga gcg gaa	192
Lys Glu Leu Gln Glu Asp Glu Leu Lys Ser Leu Gln Arg Gly Ala Glu	
50 55 60	
act act tgg gac caa agc gag gac atg gaa tgg gaa acc aca gtg gat	240
Thr Thr Trp Asp Gln Ser Glu Asp Met Glu Trp Glu Thr Thr Val Asp	
65 70 75 80	
gaa atg acc aaa aag caa gta ttc att ttt gat tct ttg gtt aaa aaa	288
Glu Met Thr Lys Lys Gln Val Phe Ile Phe Asp Ser Leu Val Lys Lys	
85 90 95	
tgt tta ttt gaa gtg ctt aac aca aag aat ata ttt cct ggt gat gtt	336
Cys Leu Phe Glu Val Leu Asn Thr Lys Asn Ile Phe Pro Gly Asp Val	
100 105 110	
aat tgg ttt gtg caa cat gaa tgg gga aaa gac caa ggc tgg cac tgc	384
Asn Trp Phe Val Gln His Glu Trp Gly Lys Asp Gln Gly Trp His Cys	
115 120 125	
cat gta cta att gga gga aag gac ttt agt caa gct caa ggg aaa tgg	432
His Val Leu Ile Gly Gly Lys Asp Phe Ser Gln Ala Gln Gly Lys Trp	
130 135 140	

tgg	aga	agg	caa	cta	aat	gtt	tac	tgg	agc	aga	tgg	ttg	gta	aca	gcc	480
Trp	Arg	Arg	Gln	Leu	Asn	Val	Tyr	Trp	Ser	Arg	Trp	Leu	Val	Thr	Ala	
145					150				155						160	
tgt	aat	gtg	caa	cta	aca	cca	gct	gaa	aga	att	aaa	cta	aga	gaa	ata	528
Cys	Asn	Val	Gln	Leu	Thr	Pro	Ala	Glu	Arg	Ile	Lys	Leu	Arg	Glu	Ile	
				165					170					175		
gca	gaa	gac	aat	gag	tgg	gtt	act	cta	ctt	act	tat	aag	cat	aag	caa	576
Ala	Glu	Asp	Asn	Glu	Trp	Val	Thr	Leu	Leu	Thr	Tyr	Lys	His	Lys	Gln	
			180					185					190			
acc	aaa	aaa	gac	tat	acc	aag	tgt	gtt	ctt	ttt	gga	aac	atg	att	gct	624
Thr	Lys	Lys	Asp	Tyr	Thr	Lys	Cys	Val	Leu	Phe	Gly	Asn	Met	Ile	Ala	
		195					200					205				
tac	tat	ttt	tta	act	aaa	aag	aaa	ata	agc	act	agt	cca	cca	aga	gac	672
Tyr	Tyr	Phe	Leu	Thr	Lys	Lys	Lys	Ile	Ser	Thr	Ser	Pro	Pro	Arg	Asp	
	210					215					220					
gga	ggc	tat	ttt	ctt	agc	agt	gac	tct	ggc	tgg	aaa	act	aac	ttt	tta	720
Gly	Gly	Tyr	Phe	Leu	Ser	Ser	Asp	Ser	Gly	Trp	Lys	Thr	Asn	Phe	Leu	
225					230					235					240	
aaa	gaa	ggc	gag	cgc	cat	cta	gtg	agc	aaa	cta	tac	act	gat	gac	atg	768
Lys	Glu	Gly	Glu	Arg	His	Leu	Val	Ser	Lys	Leu	Tyr	Thr	Asp	Asp	Met	
				245					250					255		
cgg	cca	gaa	acg	gtt	gaa	acc	aca	gta	acc	act	gcg	cag	gaa	act	aag	816
Arg	Pro	Glu	Thr	Val	Glu	Thr	Thr	Val	Thr	Thr	Ala	Gln	Glu	Thr	Lys	
			260					265					270			
cgc	ggc	aga	att	caa	act	aaa	aaa	gaa	gtt	tct	att	aaa	act	aca	ctt	864
Arg	Gly	Arg	Ile	Gln	Thr	Lys	Lys	Glu	Val	Ser	Ile	Lys	Thr	Thr	Leu	
		275					280					285				
aaa	gag	ctg	gtg	cat	aaa	aga	gta	acc	tca	cca	gag	gac	tgg	atg	atg	912
Lys	Glu	Leu	Val	His	Lys	Arg	Val	Thr	Ser	Pro	Glu	Asp	Trp	Met	Met	
		290				295					300					
atg	cag	cca	gac	agt	tac	att	gaa	atg	atg	gct	caa	cca	ggt	gga	gaa	960
Met	Gln	Pro	Asp	Ser	Tyr	Ile	Glu	Met	Met	Ala	Gln	Pro	Gly	Gly	Glu	
305					310					315					320	
aac	ctg	ctg	aaa	aat	acg	cta	gag	att	tgt	aca	cta	act	cta	gcc	aga	1008
Asn	Leu	Leu	Lys	Asn	Thr	Leu	Glu	Ile	Cys	Thr	Leu	Thr	Leu	Ala	Arg	
				325					330					335		
acc	aaa	aca	gca	ttt	gac	tta	att	tta	gaa	aaa	gct	gaa	acc	agc	aaa	1056
Thr	Lys	Thr	Ala	Phe	Asp	Leu	Ile	Leu	Glu	Lys	Ala	Glu	Thr	Ser	Lys	
			340					345					350			
cta	acc	aac	ttt	tca	ctg	cct	gac	aca	aga	acc	tgc	aga	att	ttt	gct	1104
Leu	Thr	Asn	Phe	Ser	Leu	Pro	Asp	Thr	Arg	Thr	Cys	Arg	Ile	Phe	Ala	
		355					360					365				
ttt	cat	ggc	tgg	aac	tat	gtt	aaa	gtt	tgc	cat	gct	att	tgc	tgt	gtt	1152
Phe	His	Gly	Trp	Asn	Tyr	Val	Lys	Val	Cys	His	Ala	Ile	Cys	Cys	Val	
		370				375					380					
tta	aac	aga	caa	gga	ggc	aaa	aga	aat	act	gtt	tta	ttt	cat	gga	cca	1200
Leu	Asn	Arg	Gln	Gly	Gly	Lys	Arg	Asn	Thr	Val	Leu	Phe	His	Gly	Pro	
				385		390				395					400	
gcc	agc	aca	ggc	aaa	tct	att	att	gca	caa	gcc	ata	gca	caa	gca	gtt	1248
Ala	Ser	Thr	Gly	Lys	Ser	Ile	Ile	Ala	Gln	Ala	Ile	Ala	Gln	Ala	Val	
				405					410					415		

ggc aat gtt ggt tgc tat aat gca gcc aat gta aac ttt cca ttt aat	1296
Gly Asn Val Gly Cys Tyr Asn Ala Ala Asn Val Asn Phe Pro Phe Asn	
420 425 430	
gac tgt acc aac aag aac ttg att tgg gta gaa gaa gct ggt aac ttt	1344
Asp Cys Thr Asn Lys Asn Leu Ile Trp Val Glu Glu Ala Gly Asn Phe	
435 440 445	
gga cag caa gta aac cag ttt aaa gcc att tgc tct ggt caa act att	1392
Gly Gln Gln Val Asn Gln Phe Lys Ala Ile Cys Ser Gly Gln Thr Ile	
450 455 460	
cgc att gat caa aaa gga aaa ggc agc aaa cag att gaa cca aca cca	1440
Arg Ile Asp Gln Lys Gly Lys Gly Ser Lys Gln Ile Glu Pro Thr Pro	
465 470 475 480	
gtc atc atg acc aca aat gag aac att aca gtg gtc aga ata ggc tgc	1488
Val Ile Met Thr Thr Asn Glu Asn Ile Thr Val Val Arg Ile Gly Cys	
485 490 495	
gaa gaa aga cca gaa cac act caa cca atc aga gac aga atg ctt aac	1536
Glu Glu Arg Pro Glu His Thr Gln Pro Ile Arg Asp Arg Met Leu Asn	
500 505 510	
att cat cta aca cat acc ttg cct ggt gac ttt ggt ttg gtt gac aaa	1584
Ile His Leu Thr His Thr Leu Pro Gly Asp Phe Gly Leu Val Asp Lys	
515 520 525	
aat gaa tgg ccc atg att tgt gct tgg ttg gta aag aat ggt tac caa	1632
Asn Glu Trp Pro Met Ile Cys Ala Trp Leu Val Lys Asn Gly Tyr Gln	
530 535 540	
tct acc atg gca agc tac tgt gct aaa tgg ggc aaa gtt cct gat tgg	1680
Ser Thr Met Ala Ser Tyr Cys Ala Lys Trp Gly Lys Val Pro Asp Trp	
545 550 555 560	
tca gaa aac tgg gcg gag cca aag gtg cca act cct ata aat tta cta	1728
Ser Glu Asn Trp Ala Glu Pro Lys Val Pro Thr Pro Ile Asn Leu Leu	
565 570 575	
ggc tgc gca cgc tca cca ttc acg aca ccg aaa agt acg cct ctc agc	1776
Gly Ser Ala Arg Ser Pro Phe Thr Thr Pro Lys Ser Thr Pro Leu Ser	
580 585 590	
cag aac tat gca cta act cca ctt gca tgc gat ctc gag gac ctg gct	1824
Gln Asn Tyr Ala Leu Thr Pro Leu Ala Ser Asp Leu Glu Asp Leu Ala	
595 600 605	
tta gag cct tgg agc aca cca aat act cct gtt gcg ggc act gca gaa	1872
Leu Glu Pro Trp Ser Thr Pro Asn Thr Pro Val Ala Gly Thr Ala Glu	
610 615 620	
acc cag aac act ggg gaa gct ggt tcc aaa gcc tgc caa gat ggt caa	1920
Thr Gln Asn Thr Gly Glu Ala Gly Ser Lys Ala Cys Gln Asp Gly Gln	
625 630 635 640	
ctg agc cca act tgg tca gag atc gag gag gat ttg aga gcg tgc ttc	1968
Leu Ser Pro Thr Trp Ser Glu Ile Glu Glu Asp Leu Arg Ala Cys Phe	
645 650 655	
ggc gcg gaa ccg ttg aag aaa gac ttc agc gag ccg ctg aac ttg gac	2016
Gly Ala Glu Pro Leu Lys Lys Asp Phe Ser Glu Pro Leu Asn Leu Asp	
660 665 670	
taa	2019



<210> 2  
 <211> 672  
 <212> PRT  
 <213> Wildtype Parvovirus NS1

<400> 2

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Met	Ala	Gly	Asn	Ala	Tyr	Ser	Asp	Glu	Val	Leu	Gly	Ala	Thr	Asn	Trp	1	5	10	15
Leu	Lys	Glu	Lys	Ser	Asn	Gln	Glu	Val	Phe	Ser	Phe	Val	Phe	Lys	Asn	20	25	30	
Glu	Asn	Val	Gln	Leu	Asn	Gly	Lys	Asp	Ile	Gly	Trp	Asn	Ser	Tyr	Lys	35	40	45	
Lys	Glu	Leu	Gln	Glu	Asp	Glu	Leu	Lys	Ser	Leu	Gln	Arg	Gly	Ala	Glu	50	55	60	
Thr	Thr	Trp	Asp	Gln	Ser	Glu	Asp	Met	Glu	Trp	Glu	Thr	Thr	Val	Asp	65	70	75	80
Glu	Met	Thr	Lys	Lys	Gln	Val	Phe	Ile	Phe	Asp	Ser	Leu	Val	Lys	Lys	85	90	95	
Cys	Leu	Phe	Glu	Val	Leu	Asn	Thr	Lys	Asn	Ile	Phe	Pro	Gly	Asp	Val	100	105	110	
Asn	Trp	Phe	Val	Gln	His	Glu	Trp	Gly	Lys	Asp	Gln	Gly	Trp	His	Cys	115	120	125	
His	Val	Leu	Ile	Gly	Gly	Lys	Asp	Phe	Ser	Gln	Ala	Gln	Gly	Lys	Trp	130	135	140	
Trp	Arg	Arg	Gln	Leu	Asn	Val	Tyr	Trp	Ser	Arg	Trp	Leu	Val	Thr	Ala	145	150	155	160
Cys	Asn	Val	Gln	Leu	Thr	Pro	Ala	Glu	Arg	Ile	Lys	Leu	Arg	Glu	Ile	165	170	175	
Ala	Glu	Asp	Asn	Glu	Trp	Val	Thr	Leu	Leu	Thr	Tyr	Lys	His	Lys	Gln	180	185	190	
Thr	Lys	Lys	Asp	Tyr	Thr	Lys	Cys	Val	Leu	Phe	Gly	Asn	Met	Ile	Ala	195	200	205	
Tyr	Tyr	Phe	Leu	Thr	Lys	Lys	Lys	Ile	Ser	Thr	Ser	Pro	Pro	Arg	Asp	210	215	220	
Gly	Gly	Tyr	Phe	Leu	Ser	Ser	Asp	Ser	Gly	Trp	Lys	Thr	Asn	Phe	Leu	225	230	235	240
Lys	Glu	Gly	Glu	Arg	His	Leu	Val	Ser	Lys	Leu	Tyr	Thr	Asp	Asp	Met	245	250	255	
Arg	Pro	Glu	Thr	Val	Glu	Thr	Thr	Val	Thr	Thr	Ala	Gln	Glu	Thr	Lys	260	265	270	
Arg	Gly	Arg	Ile	Gln	Thr	Lys	Lys	Glu	Val	Ser	Ile	Lys	Thr	Thr	Leu	275	280	285	
Lys	Glu	Leu	Val	His	Lys	Arg	Val	Thr	Ser	Pro	Glu	Asp	Trp	Met	Met	290	295	300	
Met	Gln	Pro	Asp	Ser	Tyr	Ile	Glu	Met	Met	Ala	Gln	Pro	Gly	Gly	Glu	305	310	315	320

Asn	Leu	Leu	Lys	Asn	Thr	Leu	Glu	Ile	Cys	Thr	Leu	Thr	Leu	Ala	Arg	
				325					330					335		
Thr	Lys	Thr	Ala	Phe	Asp	Leu	Ile	Leu	Glu	Lys	Ala	Glu	Thr	Ser	Lys	
			340					345					350			
Leu	Thr	Asn	Phe	Ser	Leu	Pro	Asp	Thr	Arg	Thr	Cys	Arg	Ile	Phe	Ala	
		355					360					365				
Phe	His	Gly	Trp	Asn	Tyr	Val	Lys	Val	Cys	His	Ala	Ile	Cys	Cys	Val	
	370					375					380					
Leu	Asn	Arg	Gln	Gly	Gly	Lys	Arg	Asn	Thr	Val	Leu	Phe	His	Gly	Pro	
385					390					395					400	
Ala	Ser	Thr	Gly	Lys	Ser	Ile	Ile	Ala	Gln	Ala	Ile	Ala	Gln	Ala	Val	
				405					410					415		
Gly	Asn	Val	Gly	Cys	Tyr	Asn	Ala	Ala	Asn	Val	Asn	Phe	Pro	Phe	Asn	
			420					425					430			
Asp	Cys	Thr	Asn	Lys	Asn	Leu	Ile	Trp	Val	Glu	Glu	Ala	Gly	Asn	Phe	
			435				440					445				
Gly	Gln	Gln	Val	Asn	Gln	Phe	Lys	Ala	Ile	Cys	Ser	Gly	Gln	Thr	Ile	
	450					455					460					
Arg	Ile	Asp	Gln	Lys	Gly	Lys	Gly	Ser	Lys	Gln	Ile	Glu	Pro	Thr	Pro	
465					470					475					480	
Val	Ile	Met	Thr	Thr	Asn	Glu	Asn	Ile	Thr	Val	Val	Arg	Ile	Gly	Cys	
				485					490					495		
Glu	Glu	Arg	Pro	Glu	His	Thr	Gln	Pro	Ile	Arg	Asp	Arg	Met	Leu	Asn	
			500					505					510			
Ile	His	Leu	Thr	His	Thr	Leu	Pro	Gly	Asp	Phe	Gly	Leu	Val	Asp	Lys	
		515					520					525				
Asn	Glu	Trp	Pro	Met	Ile	Cys	Ala	Trp	Leu	Val	Lys	Asn	Gly	Tyr	Gln	
	530					535					540					
Ser	Thr	Met	Ala	Ser	Tyr	Cys	Ala	Lys	Trp	Gly	Lys	Val	Pro	Asp	Trp	
545					550					555					560	
Ser	Glu	Asn	Trp	Ala	Glu	Pro	Lys	Val	Pro	Thr	Pro	Ile	Asn	Leu	Leu	
				565					570					575		
Gly	Ser	Ala	Arg	Ser	Pro	Phe	Thr	Thr	Pro	Lys	Ser	Thr	Pro	Leu	Ser	
			580					585					590			
Gln	Asn	Tyr	Ala	Leu	Thr	Pro	Leu	Ala	Ser	Asp	Leu	Glu	Asp	Leu	Ala	
		595					600					605				
Leu	Glu	Pro	Trp	Ser	Thr	Pro	Asn	Thr	Pro	Val	Ala	Gly	Thr	Ala	Glu	
	610					615					620					
Thr	Gln	Asn	Thr	Gly	Glu	Ala	Gly	Ser	Lys	Ala	Cys	Gln	Asp	Gly	Gln	
625					630					635					640	
Leu	Ser	Pro	Thr	Trp	Ser	Glu	Ile	Glu	Glu	Asp	Leu	Arg	Ala	Cys	Phe	
				645					650					655		
Gly	Ala	Glu	Pro	Leu	Lys	Lys	Asp	Phe	Ser	Glu	Pro	Leu	Asn	Leu	Asp	
			660					665					670			

<210> 3  
 <211> 60  
 <212> DNA  
 <213> part of Parvovirus NS1 variant

<220>  
 <221> CDS  
 <222> (1)..(60)

<400> 3

gaa gtt gct att aaa act aca ctt aaa gag ctg gtg cat aaa aga gta 48  
 Glu Val Ala Ile Lys Thr Thr Leu Lys Glu Leu Val His Lys Arg Val  
 1 5 10 15

acc tca cca gag 60  
 Thr Ser Pro Glu  
 20

<210> 4  
 <211> 2019  
 <212> DNA  
 <213> Parvovirus NS1 variant

<220>  
 <221> CDS  
 <222> (1)..(2016)

<400> 4

atg gct gga aat gct tac tct gat gaa gtt ttg gga gca acc aac tgg 48  
 Met Ala Gly Asn Ala Tyr Ser Asp Glu Val Leu Gly Ala Thr Asn Trp  
 1 5 10 15

tta aag gaa aaa agt aac cag gaa gtg ttc tca ttt gtt ttt aaa aat 96  
 Leu Lys Glu Lys Ser Asn Gln Glu Val Phe Ser Phe Val Phe Lys Asn  
 20 25 30

gaa aat gtt caa ctg aat gga aaa gat atc gga tgg aat agt tac aaa 144  
 Glu Asn Val Gln Leu Asn Gly Lys Asp Ile Gly Trp Asn Ser Tyr Lys  
 35 40 45

aaa gag ctg cag gag gac gag ctg aaa tct tta caa cga gga gcg gaa 192  
 Lys Glu Leu Gln Glu Asp Glu Leu Lys Ser Leu Gln Arg Gly Ala Glu  
 50 55 60

act act tgg gac caa agc gag gac atg gaa tgg gaa acc aca gtg gat 240  
 Thr Thr Trp Asp Gln Ser Glu Asp Met Glu Trp Glu Thr Thr Val Asp  
 65 70 75 80

gaa atg acc aaa aag caa gta ttc att ttt gat tct ttg gtt aaa aaa 288  
 Glu Met Thr Lys Lys Gln Val Phe Ile Phe Asp Ser Leu Val Lys Lys  
 85 90 95

tgt tta ttt gaa gtg ctt aac aca aag aat ata ttt cct ggt gat gtt 336  
 Cys Leu Phe Glu Val Leu Asn Thr Lys Asn Ile Phe Pro Gly Asp Val  
 100 105 110

aat tgg ttt gtg caa cat gaa tgg gga aaa gac caa ggc tgg cac tgc	384
Asn Trp Phe Val Gln His Glu Trp Gly Lys Asp Gln Gly Trp His Cys	
115 120 125	
cat gta cta att gga gga aag gac ttt agt caa gct caa ggg aaa tgg	432
His Val Leu Ile Gly Gly Lys Asp Phe Ser Gln Ala Gln Gly Lys Trp	
130 135 140	
tgg aga agg caa cta aat gtt tac tgg agc aga tgg ttg gta aca gcc	480
Trp Arg Arg Gln Leu Asn Val Tyr Trp Ser Arg Trp Leu Val Thr Ala	
145 150 155 160	
tgt aat gtg caa cta aca cca gct gaa aga att aaa cta aga gaa ata	528
Cys Asn Val Gln Leu Thr Pro Ala Glu Arg Ile Lys Leu Arg Glu Ile	
165 170 175	
gca gaa gac aat gag tgg gtt act cta ctt act tat aag cat aag caa	576
Ala Glu Asp Asn Glu Trp Val Thr Leu Leu Thr Tyr Lys His Lys Gln	
180 185 190	
acc aaa aaa gac tat acc aag tgt gtt ctt ttt gga aac atg att gct	624
Thr Lys Lys Asp Tyr Thr Lys Cys Val Leu Phe Gly Asn Met Ile Ala	
195 200 205	
tac tat ttt tta act aaa aag aaa ata agc act agt cca cca aga gac	672
Tyr Tyr Phe Leu Thr Lys Lys Lys Ile Ser Thr Ser Pro Pro Arg Asp	
210 215 220	
gga ggc tat ttt ctt agc agt gac tct ggc tgg aaa act aac ttt tta	720
Gly Gly Tyr Phe Leu Ser Ser Asp Ser Gly Trp Lys Thr Asn Phe Leu	
225 230 235 240	
aaa gaa ggc gag cgc cat cta gtg agc aaa cta tac act gat gac atg	768
Lys Glu Gly Glu Arg His Leu Val Ser Lys Leu Tyr Thr Asp Asp Met	
245 250 255	
cgg cca gaa acg gtt gaa acc aca gta acc act gcg cag gaa act aag	816
Arg Pro Glu Thr Val Glu Thr Thr Val Thr Thr Ala Gln Glu Thr Lys	
260 265 270	
cgc ggc aga att caa act aaa aaa gaa gtt gct att aaa act aca ctt	864
Arg Gly Arg Ile Gln Thr Lys Lys Glu Val Ala Ile Lys Thr Thr Leu	
275 280 285	
aaa gag ctg gtg cat aaa aga gta acc tca cca gag gac tgg atg atg	912
Lys Glu Leu Val His Lys Arg Val Thr Ser Pro Glu Asp Trp Met Met	
290 295 300	
atg cag cca gac agt tac att gaa atg atg gct caa cca ggt gga gaa	960
Met Gln Pro Asp Ser Tyr Ile Glu Met Met Ala Gln Pro Gly Gly Glu	
305 310 315 320	
aac ctg ctg aaa aat acg cta gag att tgt aca cta act cta gcc aga	1008
Asn Leu Leu Lys Asn Thr Leu Glu Ile Cys Thr Leu Thr Leu Ala Arg	
325 330 335	
acc aaa aca gca ttt gac tta att tta gaa aaa gct gaa acc agc aaa	1056
Thr Lys Thr Ala Phe Asp Leu Ile Leu Glu Lys Ala Glu Thr Ser Lys	
340 345 350	
cta acc aac ttt tca ctg cct gac aca aga acc tgc aga att ttt gct	1104
Leu Thr Asn Phe Ser Leu Pro Asp Thr Arg Thr Cys Arg Ile Phe Ala	
355 360 365	
ttt cat ggc tgg aac tat gtt aaa gtt tgc cat gct att tgc tgt gtt	1152
Phe His Gly Trp Asn Tyr Val Lys Val Cys His Ala Ile Cys Cys Val	
370 375 380	

tta aac aga caa gga ggc aaa aga aat act gtt tta ttt cat gga cca	1200
Leu Asn Arg Gln Gly Gly Lys Arg Asn Thr Val Leu Phe His Gly Pro	
385 390 395 400	
gcc agc aca ggc aaa tct att att gca caa gcc ata gca caa gca gtt	1248
Ala Ser Thr Gly Lys Ser Ile Ile Ala Gln Ala Ile Ala Gln Ala Val	
405 410 415	
ggc aat gtt ggt tgc tat aat gca gcc aat gta aac ttt cca ttt aat	1296
Gly Asn Val Gly Cys Tyr Asn Ala Ala Asn Val Asn Phe Pro Phe Asn	
420 425 430	
gac tgt acc aac aag aac ttg att tgg gta gaa gaa gct ggt aac ttt	1344
Asp Cys Thr Asn Lys Asn Leu Ile Trp Val Glu Glu Ala Gly Asn Phe	
435 440 445	
gga cag caa gta aac cag ttt aaa gcc att tgc tct ggt caa act att	1392
Gly Gln Gln Val Asn Gln Phe Lys Ala Ile Cys Ser Gly Gln Thr Ile	
450 455 460	
cgc att gat caa aaa gga aaa ggc agc aaa cag att gaa cca aca cca	1440
Arg Ile Asp Gln Lys Gly Lys Gly Ser Lys Gln Ile Glu Pro Thr Pro	
465 470 475 480	
gtc atc atg acc aca aat gag aac att aca gtg gtc aga ata ggc tgc	1488
Val Ile Met Thr Thr Asn Glu Asn Ile Thr Val Val Arg Ile Gly Cys	
485 490 495	
gaa gaa aga cca gaa cac act caa cca atc aga gac aga atg ctt aac	1536
Glu Glu Arg Pro Glu His Thr Gln Pro Ile Arg Asp Arg Met Leu Asn	
500 505 510	
att cat cta aca cat acc ttg cct ggt gac ttt ggt ttg gtt gac aaa	1584
Ile His Leu Thr His Thr Leu Pro Gly Asp Phe Gly Leu Val Asp Lys	
515 520 525	
aat gaa tgg ccc atg att tgt gct tgg ttg gta aag aat ggt tac caa	1632
Asn Glu Trp Pro Met Ile Cys Ala Trp Leu Val Lys Asn Gly Tyr Gln	
530 535 540	
tct acc atg gca agc tac tgt gct aaa tgg ggc aaa gtt cct gat tgg	1680
Ser Thr Met Ala Ser Tyr Cys Ala Lys Trp Gly Lys Val Pro Asp Trp	
545 550 555 560	
tca gaa aac tgg gcg gag cca aag gtg cca act cct ata aat tta cta	1728
Ser Glu Asn Trp Ala Glu Pro Lys Val Pro Thr Pro Ile Asn Leu Leu	
565 570 575	
ggc tgc gca cgc tca cca ttc acg aca ccg aaa agt acg cct ctc agc	1776
Gly Ser Ala Arg Ser Pro Phe Thr Thr Pro Lys Ser Thr Pro Leu Ser	
580 585 590	
cag aac tat gca cta act cca ctt gca tgc gat ctc gag gac ctg gct	1824
Gln Asn Tyr Ala Leu Thr Pro Leu Ala Ser Asp Leu Glu Asp Leu Ala	
595 600 605	
tta gag cct tgg agc aca cca aat act cct gtt gcg ggc act gca gaa	1872
Leu Glu Pro Trp Ser Thr Pro Asn Thr Pro Val Ala Gly Thr Ala Glu	
610 615 620	
acc cag aac act ggg gaa gct ggt tcc aaa gcc tgc caa gat ggt caa	1920
Thr Gln Asn Thr Gly Glu Ala Gly Ser Lys Ala Cys Gln Asp Gly Gln	
625 630 635 640	
ctg agc cca act tgg tca gag atc gag gag gat ttg aga gcg tgc ttc	1968
Leu Ser Pro Thr Trp Ser Glu Ile Glu Glu Asp Leu Arg Ala Cys Phe	
645 650 655	

ggt gcg gaa ccg ttg aag aaa gac ttc agc gag ccg ctg aac ttg gac 2016  
 Gly Ala Glu Pro Leu Lys Lys Asp Phe Ser Glu Pro Leu Asn Leu Asp  
                   660                  665                  670

taa 2019

<210> 5  
 <211> 20  
 <212> PRT  
 <213> part of Parvovirus NS1 variant

<400> 5

Glu Val Ala Ile Lys Thr Thr Leu Lys Glu Leu Val His Lys Arg Val  
 1                  5                  10                  15  
 Thr Ser Pro Glu  
                   20

<210> 6  
 <211> 672  
 <212> PRT  
 <213> Parvovirus NS1 variant

<400> 6

Met Ala Gly Asn Ala Tyr Ser Asp Glu Val Leu Gly Ala Thr Asn Trp  
 1                  5                  10                  15  
 Leu Lys Glu Lys Ser Asn Gln Glu Val Phe Ser Phe Val Phe Lys Asn  
                   20                  25                  30  
 Glu Asn Val Gln Leu Asn Gly Lys Asp Ile Gly Trp Asn Ser Tyr Lys  
                   35                  40                  45  
 Lys Glu Leu Gln Glu Asp Glu Leu Lys Ser Leu Gln Arg Gly Ala Glu  
                   50                  55                  60  
 Thr Thr Trp Asp Gln Ser Glu Asp Met Glu Trp Glu Thr Thr Val Asp  
 65                  70                  75                  80  
 Glu Met Thr Lys Lys Gln Val Phe Ile Phe Asp Ser Leu Val Lys Lys  
                   85                  90                  95  
 Cys Leu Phe Glu Val Leu Asn Thr Lys Asn Ile Phe Pro Gly Asp Val  
                   100                  105                  110  
 Asn Trp Phe Val Gln His Glu Trp Gly Lys Asp Gln Gly Trp His Cys  
 115                  120                  125  
 His Val Leu Ile Gly Gly Lys Asp Phe Ser Gln Ala Gln Gly Lys Trp  
 130                  135                  140  
 Trp Arg Arg Gln Leu Asn Val Tyr Trp Ser Arg Trp Leu Val Thr Ala  
 145                  150                  155                  160  
 Cys Asn Val Gln Leu Thr Pro Ala Glu Arg Ile Lys Leu Arg Glu Ile  
                   165                  170                  175  
 Ala Glu Asp Asn Glu Trp Val Thr Leu Leu Thr Tyr Lys His Lys Gln  
                   180                  185                  190

10

Thr	Lys	Lys	Asp	Tyr	Thr	Lys	Cys	Val	Leu	Phe	Gly	Asn	Met	Ile	Ala	
	195						200					205				
Tyr	Tyr	Phe	Leu	Thr	Lys	Lys	Lys	Ile	Ser	Thr	Ser	Pro	Pro	Arg	Asp	
	210					215					220					
Gly	Gly	Tyr	Phe	Leu	Ser	Ser	Asp	Ser	Gly	Trp	Lys	Thr	Asn	Phe	Leu	
<del>225</del>					<del>230</del>					<del>235</del>					<del>240</del>	
Lys	Glu	Gly	Glu	Arg	His	Leu	Val	Ser	Lys	Leu	Tyr	Thr	Asp	Asp	Met	
				245					250					255		
Arg	Pro	Glu	Thr	Val	Glu	Thr	Thr	Val	Thr	Thr	Ala	Gln	Glu	Thr	Lys	
			260					265					270			
Arg	Gly	Arg	Ile	Gln	Thr	Lys	Lys	Glu	Val	Ala	Ile	Lys	Thr	Thr	Leu	
		275					280					285				
Lys	Glu	Leu	Val	His	Lys	Arg	Val	Thr	Ser	Pro	Glu	Asp	Trp	Met	Met	
	290					295					300					
Met	Gln	Pro	Asp	Ser	Tyr	Ile	Glu	Met	Met	Ala	Gln	Pro	Gly	Gly	Glu	
305					310					315					320	
Asn	Leu	Leu	Lys	Asn	Thr	Leu	Glu	Ile	Cys	Thr	Leu	Thr	Leu	Ala	Arg	
				325					330					335		
Thr	Lys	Thr	Ala	Phe	Asp	Leu	Ile	Leu	Glu	Lys	Ala	Glu	Thr	Ser	Lys	
			340					345					350			
Leu	Thr	Asn	Phe	Ser	Leu	Pro	Asp	Thr	Arg	Thr	Cys	Arg	Ile	Phe	Ala	
		355					360					365				
Phe	His	Gly	Trp	Asn	Tyr	Val	Lys	Val	Cys	His	Ala	Ile	Cys	Cys	Val	
	370					375					380					
Leu	Asn	Arg	Gln	Gly	Gly	Lys	Arg	Asn	Thr	Val	Leu	Phe	His	Gly	Pro	
385					390					395					400	
Ala	Ser	Thr	Gly	Lys	Ser	Ile	Ile	Ala	Gln	Ala	Ile	Ala	Gln	Ala	Val	
				405					410					415		
Gly	Asn	Val	Gly	Cys	Tyr	Asn	Ala	Ala	Asn	Val	Asn	Phe	Pro	Phe	Asn	
			420					425					430			
Asp	Cys	Thr	Asn	Lys	Asn	Leu	Ile	Trp	Val	Glu	Glu	Ala	Gly	Asn	Phe	
		435				440						445				
Gly	Gln	Gln	Val	Asn	Gln	Phe	Lys	Ala	Ile	Cys	Ser	Gly	Gln	Thr	Ile	
	450					455					460					
Arg	Ile	Asp	Gln	Lys	Gly	Lys	Gly	Ser	Lys	Gln	Ile	Glu	Pro	Thr	Pro	
465					470					475					480	
Val	Ile	Met	Thr	Thr	Asn	Glu	Asn	Ile	Thr	Val	Val	Arg	Ile	Gly	Cys	
				485					490					495		
Glu	Glu	Arg	Pro	Glu	His	Thr	Gln	Pro	Ile	Arg	Asp	Arg	Met	Leu	Asn	
			500					505					510			
Ile	His	Leu	Thr	His	Thr	Leu	Pro	Gly	Asp	Phe	Gly	Leu	Val	Asp	Lys	
		515					520					525				
Asn	Glu	Trp	Pro	Met	Ile	Cys	Ala	Trp	Leu	Val	Lys	Asn	Gly	Tyr	Gln	
	530					535					540					

11

Ser Thr Met Ala Ser Tyr Cys Ala Lys Trp Gly Lys Val Pro Asp Trp  
545 550 555 560

Ser Glu Asn Trp Ala Glu Pro Lys Val Pro Thr Pro Ile Asn Leu Leu  
565 570 575

Gly Ser Ala Arg Ser Pro Phe Thr Thr Pro Lys Ser Thr Pro Leu Ser  
580 585 590

Gln Asn Tyr Ala Leu Thr Pro Leu Ala Ser Asp Leu Glu Asp Leu Ala  
595 600 605

Leu Glu Pro Trp Ser Thr Pro Asn Thr Pro Val Ala Gly Thr Ala Glu  
610 615 620

Thr Gln Asn Thr Gly Glu Ala Gly Ser Lys Ala Cys Gln Asp Gly Gln  
625 630 635 640

Leu Ser Pro Thr Trp Ser Glu Ile Glu Glu Asp Leu Arg Ala Cys Phe  
645 650 655

Gly Ala Glu Pro Leu Lys Lys Asp Phe Ser Glu Pro Leu Asn Leu Asp  
660 665 670

&lt;210&gt; 7

&lt;211&gt; 60

&lt;212&gt; DNA

&lt;213&gt; part of Parvovirus NS1 variant

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (1)..(60)

&lt;400&gt; 7

aca aga gcc tgc aga att ttt gct ttt cat ggc tgg aac tat gtt aaa 48  
Thr Arg Ala Cys Arg Ile Phe Ala Phe His Gly Trp Asn Tyr Val Lys  
1 5 10 15

gtt tgc cat gct 60  
Val Cys His Ala  
20

&lt;210&gt; 8

&lt;211&gt; 2019

&lt;212&gt; DNA

&lt;213&gt; Parvovirus NS1 variant

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (1)..(2016)

&lt;400&gt; 8

atg gct gga aat gct tac tct gat gaa gtt ttg gga gca acc aac tgg 48  
Met Ala Gly Asn Ala Tyr Ser Asp Glu Val Leu Gly Ala Thr Asn Trp  
1 5 10 15

tta aag gaa aaa agt aac cag gaa gtg ttc tca ttt gtt ttt aaa aat 96  
Leu Lys Glu Lys Ser Asn Gln Glu Val Phe Ser Phe Val Phe Lys Asn  
20 25 30



12

gaa aat gtt caa ctg aat gga aaa gat atc gga tgg aat agt tac aaa Glu Asn Val Gln Leu Asn Gly Lys Asp Ile Gly Trp Asn Ser Tyr Lys 35 40 45	144
aaa gag ctg cag gag gac gag ctg aaa tct tta caa cga gga gcg gaa Lys Glu Leu Gln Glu Asp Ser Leu Lys Ser Leu Gln Arg Gly Ala Glu 50 55 60	192
act act tgg gac caa agc gag gac atg gaa tgg gaa acc aca gtg gat Thr Thr Trp Asp Gln Ser Glu Asp Met Glu Trp Glu Thr Thr Val Asp 65 70 75 80	240
gaa atg acc aaa aag caa gta ttc att ttt gat tct ttg gtt aaa aaa Glu Met Thr Lys Lys Gln Val Phe Ile Phe Asp Ser Leu Val Lys Lys 85 90 95	288
tgt tta ttt gaa gtg ctt aac aca aag aat ata ttt cct ggt gat gtt Cys Leu Phe Glu Val Leu Asn Thr Lys Asn Ile Phe Pro Gly Asp Val 100 105 110	336
aat tgg ttt gtg caa cat gaa tgg gga aaa gac caa ggc tgg cac tgc Asn Trp Phe Val Gln His Glu Trp Gly Lys Asp Gln Gly Trp His Cys 115 120 125	384
cat gta cta att gga gga aag gac ttt agt caa gct caa ggg aaa tgg His Val Leu Ile Gly Gly Lys Asp Phe Ser Gln Ala Gln Gly Lys Trp 130 135 140	432
tgg aga agg caa cta aat gtt tac tgg agc aga tgg ttg gta aca gcc Trp Arg Arg Gln Leu Asn Val Tyr Trp Ser Arg Trp Leu Val Thr Ala 145 150 155 160	480
tgt aat gtg caa cta aca cca gct gaa aga att aaa cta aga gaa ata Cys Asn Val Gln Leu Thr Pro Ala Glu Arg Ile Lys Leu Arg Glu Ile 165 170 175	528
gca gaa gac aat gag tgg gtt act cta ctt act tat aag cat aag caa Ala Glu Asp Asn Glu Trp Val Thr Leu Leu Thr Tyr Lys His Lys Gln 180 185 190	576
acc aaa aaa gac tat acc aag tgt gtt ctt ttt gga aac atg att gct Thr Lys Lys Asp Tyr Thr Lys Cys Val Leu Phe Gly Asn Met Ile Ala 195 200 205	624
tac tat ttt tta act aaa aag aaa ata agc act agt cca cca aga gac Tyr Tyr Phe Leu Thr Lys Lys Lys Ile Ser Thr Ser Pro Pro Arg Asp 210 215 220	672
gga ggc tat ttt ctt agc agt gac tct ggc tgg aaa act aac ttt tta Gly Gly Tyr Phe Leu Ser Ser Asp Ser Gly Trp Lys Thr Asn Phe Leu 225 230 235 240	720
aaa gaa ggc gag cgc cat cta gtg agc aaa cta tac act gat gac atg Lys Glu Gly Glu Arg His Leu Val Ser Lys Leu Tyr Thr Asp Asp Met 245 250 255	768
cgg cca gaa acg gtt gaa acc aca gta acc act gcg cag gaa act aag Arg Pro Glu Thr Val Glu Thr Thr Val Thr Thr Ala Gln Glu Thr Lys 260 265 270	816
cgc ggc aga att caa act aaa aaa gaa gtt tct att aaa act aca ctt Arg Gly Arg Ile Gln Thr Lys Lys Glu Val Ser Ile Lys Thr Thr Leu 275 280 285	864

aaa gag ctg gtg cat aaa aga gta acc tca cca gag gac tgg atg atg	912
Lys Glu Leu Val His Lys Arg Val Thr Ser Pro Glu Asp Trp Met Met	
290 295 300	
atg cag cca gac agt tac att gaa atg atg gct caa cca ggt gga gaa	960
Met Gln Pro Asp Ser Tyr Ile Glu Met Met Ala Gln Pro Gly Gly Glu	
305 310 315 320	
aac ctg ctg aaa aat acg cta gag att tgt aca cta act cta gcc aga	1008
Asn Leu Leu Lys Asn Thr Leu Glu Ile Cys Thr Leu Thr Leu Ala Arg	
325 330 335	
acc aaa aca gca ttt gac tta att tta gaa aaa gct gaa acc agc aaa	1056
Thr Lys Thr Ala Phe Asp Leu Ile Leu Glu Lys Ala Glu Thr Ser Lys	
340 345 350	
cta acc aac ttt tca ctg cct gac aca aga gcc tgc aga att ttt gct	1104
Leu Thr Asn Phe Ser Leu Pro Asp Thr Arg Ala Cys Arg Ile Phe Ala	
355 360 365	
ttt cat ggc tgg aac tat gtt aaa gtt tgc cat gct att tgc tgt gtt	1152
Phe His Gly Trp Asn Tyr Val Lys Val Cys His Ala Ile Cys Cys Val	
370 375 380	
tta aac aga caa gga ggc aaa aga aat act gtt tta ttt cat gga cca	1200
Leu Asn Arg Gln Gly Gly Lys Arg Asn Thr Val Leu Phe His Gly Pro	
385 390 395 400	
gcc agc aca ggc aaa tct att att gca caa gcc ata gca caa gca gtt	1248
Ala Ser Thr Gly Lys Ser Ile Ile Ala Gln Ala Ile Ala Gln Ala Val	
405 410 415	
ggc aat gtt ggt tgc tat aat gca gcc aat gta aac ttt cca ttt aat	1296
Gly Asn Val Gly Cys Tyr Asn Ala Ala Asn Val Asn Phe Pro Phe Asn	
420 425 430	
gac tgt acc aac aag aac ttg att tgg gta gaa gaa gct ggt aac ttt	1344
Asp Cys Thr Asn Lys Asn Leu Ile Trp Val Glu Glu Gly Asn Phe	
435 440 445	
gga cag caa gta aac cag ttt aaa gcc att tgc tct ggt caa act att	1392
Gly Gln Gln Val Asn Gln Phe Lys Ala Ile Cys Ser Gly Gln Thr Ile	
450 455 460	
cgc att gat caa aaa gga aaa ggc agc aaa cag att gaa cca aca cca	1440
Arg Ile Asp Gln Lys Gly Lys Gly Ser Lys Gln Ile Glu Pro Thr Pro	
465 470 475 480	
gtc atc atg acc aca aat gag aac att aca gtg gtc aga ata ggc tgc	1488
Val Ile Met Thr Thr Asn Glu Asn Ile Thr Val Val Arg Ile Gly Cys	
485 490 495	
gaa gaa aga cca gaa cac act caa cca atc aga gac aga atg ctt aac	1536
Glu Glu Arg Pro Glu His Thr Gln Pro Ile Arg Asp Arg Met Leu Asn	
500 505 510	
att cat cta aca cat acc ttg cct ggt gac ttt ggt ttg gtt gac aaa	1584
Ile His Leu Thr His Thr Leu Pro Gly Asp Phe Gly Leu Val Asp Lys	
515 520 525	
aat gaa tgg ccc atg att tgt gct tgg ttg gta aag aat ggt tac caa	1632
Asn Glu Trp Pro Met Ile Cys Ala Trp Leu Val Lys Asn Gly Tyr Gln	
530 535 540	
tct acc atg gca agc tac tgt gct aaa tgg ggc aaa gtt cct gat tgg	1680
Ser Thr Met Ala Ser Tyr Cys Ala Lys Trp Gly Lys Val Pro Asp Trp	
545 550 555 560	

14

tca gaa aac tgg gcg gag cca aag gtg cca act cct ata aat tta cta 1728  
 Ser Glu Asn Trp Ala Glu Pro Lys Val Pro Thr Pro Ile Asn Leu Leu  
 565 570 575

ggg tcg gca cgc tca cca ttc acg aca ccg aaa agt acg cct ctc agc 1776  
 Gly Ser Ala Arg Ser Pro Phe Thr Thr Pro Lys Ser Thr Pro Leu Ser  
 580 585 590

cag aac tat gca cta act cca ctt gca tcg gat ctc gag gac ctg gct 1824  
 Gln Asn Tyr Ala Leu Thr Pro Leu Ala Ser Asp Leu Glu Asp Leu Ala  
 595 600 605

tta gag cct tgg agc aca cca aat act cct gtt gcg ggc act gca gaa 1872  
 Leu Glu Pro Trp Ser Thr Pro Asn Thr Pro Val Ala Gly Thr Ala Glu  
 610 615 620

acc cag aac act ggg gaa gct ggt tcc aaa gcc tgc caa gat ggt caa 1920  
 Thr Gln Asn Thr Gly Glu Ala Gly Ser Lys Ala Cys Gln Asp Gly Gln  
 625 630 635 640

ctg agc cca act tgg tca gag atc gag gag gat ttg aga gcg tgc ttc 1968  
 Leu Ser Pro Thr Trp Ser Glu Ile Glu Glu Asp Leu Arg Ala Cys Phe  
 645 650 655

ggg gcg gaa ccg ttg aag aaa gac ttc agc gag ccg ctg aac ttg gac 2016  
 Gly Ala Glu Pro Leu Lys Lys Asp Phe Ser Glu Pro Leu Asn Leu Asp  
 660 665 670

taa 2019

<210> 9

<211> 20

<212> PRT

<213> part of Parvovirus NS1 variant

<400> 9

Thr Arg Ala Cys Arg Ile Phe Ala Phe His Gly Trp Asn Tyr Val Lys  
 1 5 10 15

Val Cys His Ala  
 20

<210> 10

<211> 672

<212> PRT

<213> Parvovirus NS1 variant

<400> 10

Met Ala Gly Asn Ala Tyr Ser Asp Glu Val Leu Gly Ala Thr Asn Trp  
 1 5 10 15

Leu Lys Glu Lys Ser Asn Gln Glu Val Phe Ser Phe Val Phe Lys Asn  
 20 25 30

Glu Asn Val Gln Leu Asn Gly Lys Asp Ile Gly Trp Asn Ser Tyr Lys  
 35 40 45

Lys Glu Leu Gln Glu Asp Glu Leu Lys Ser Leu Gln Arg Gly Ala Glu  
 50 55 60

15

Thr 65	Thr	Trp	Asp	Gln	Ser 70	Glu	Asp	Met	Glu	Trp 75	Glu	Thr	Thr	Val	Asp 80
Glu	Met	Thr	Lys	Lys 85	Gln	Val	Phe	Ile	Phe 90	Asp	Ser	Leu	Val	Lys 95	Lys
Cys	Leu	Phe	Glu	Val	Leu	Asn	Thr	Lys	Asn	Ile	Phe	Pro	Gly	Asp	Val
			100					105					110		
Asn	Trp	Phe	Val	Gln	His	Glu	Trp	Gly	Lys	Asp	Gln	Gly	Trp	His	Cys
		115					120					125			
His	Val	Leu	Ile	Gly	Gly	Lys	Asp	Phe	Ser	Gln	Ala	Gln	Gly	Lys	Trp
	130					135					140				
Trp	Arg	Arg	Gln	Leu	Asn	Val	Tyr	Trp	Ser	Arg	Trp	Leu	Val	Thr	Ala
145					150					155					160
Cys	Asn	Val	Gln	Leu	Thr	Pro	Ala	Glu	Arg	Ile	Lys	Leu	Arg	Glu	Ile
				165					170					175	
Ala	Glu	Asp	Asn	Glu	Trp	Val	Thr	Leu	Leu	Thr	Tyr	Lys	His	Lys	Gln
			180					185					190		
Thr	Lys	Lys	Asp	Tyr	Thr	Lys	Cys	Val	Leu	Phe	Gly	Asn	Met	Ile	Ala
		195					200					205			
Tyr	Tyr	Phe	Leu	Thr	Lys	Lys	Lys	Ile	Ser	Thr	Ser	Pro	Pro	Arg	Asp
	210					215					220				
Gly	Gly	Tyr	Phe	Leu	Ser	Ser	Asp	Ser	Gly	Trp	Lys	Thr	Asn	Phe	Leu
225					230					235					240
Lys	Glu	Gly	Glu	Arg	His	Leu	Val	Ser	Lys	Leu	Tyr	Thr	Asp	Asp	Met
				245					250					255	
Arg	Pro	Glu	Thr	Val	Glu	Thr	Thr	Val	Thr	Thr	Ala	Gln	Glu	Thr	Lys
			260					265					270		
Arg	Gly	Arg	Ile	Gln	Thr	Lys	Lys	Glu	Val	Ser	Ile	Lys	Thr	Thr	Leu
		275					280					285			
Lys	Glu	Leu	Val	His	Lys	Arg	Val	Thr	Ser	Pro	Glu	Asp	Trp	Met	Met
	290					295					300				
Met	Gln	Pro	Asp	Ser	Tyr	Ile	Glu	Met	Met	Ala	Gln	Pro	Gly	Gly	Glu
305					310					315					320
Asn	Leu	Leu	Lys	Asn	Thr	Leu	Glu	Ile	Cys	Thr	Leu	Thr	Leu	Ala	Arg
				325					330					335	
Thr	Lys	Thr	Ala	Phe	Asp	Leu	Ile	Leu	Glu	Lys	Ala	Glu	Thr	Ser	Lys
			340					345					350		
Leu	Thr	Asn	Phe	Ser	Leu	Pro	Asp	Thr	Arg	Ala	Cys	Arg	Ile	Phe	Ala
		355					360					365			
Phe	His	Gly	Trp	Asn	Tyr	Val	Lys	Val	Cys	His	Ala	Ile	Cys	Cys	Val
	370					375					380				
Leu	Asn	Arg	Gln	Gly	Gly	Lys	Arg	Asn	Thr	Val	Leu	Phe	His	Gly	Pro
385					390					395					400
Ala	Ser	Thr	Gly	Lys	Ser	Ile	Ile	Ala	Gln	Ala	Ile	Ala	Gln	Ala	Val
				405					410					415	

16

Gly Asn Val Gly Cys Tyr Asn Ala Ala Asn Val Asn Phe Pro Phe Asn  
 420 425 430  
 Asp Cys Thr Asn Lys Asn Leu Ile Trp Val Glu Glu Ala Gly Asn Phe  
 435 440 445  
 Gly Gln Gln Val Asn Gln Phe Lys Ala Ile Cys Ser Gly Gln Thr Ile  
 450 455 460  
 Arg Ile Asp Gln Lys Gly Lys Gly Ser Lys Gln Ile Glu Pro Thr Pro  
 465 470 475 480  
 Val Ile Met Thr Thr Asn Glu Asn Ile Thr Val Val Arg Ile Gly Cys  
 485 490 495  
 Glu Glu Arg Pro Glu His Thr Gln Pro Ile Arg Asp Arg Met Leu Asn  
 500 505 510  
 Ile His Leu Thr His Thr Leu Pro Gly Asp Phe Gly Leu Val Asp Lys  
 515 520 525  
 Asn Glu Trp Pro Met Ile Cys Ala Trp Leu Val Lys Asn Gly Tyr Gln  
 530 535 540  
 Ser Thr Met Ala Ser Tyr Cys Ala Lys Trp Gly Lys Val Pro Asp Trp  
 545 550 555 560  
 Ser Glu Asn Trp Ala Glu Pro Lys Val Pro Thr Pro Ile Asn Leu Leu  
 565 570 575  
 Gly Ser Ala Arg Ser Pro Phe Thr Thr Pro Lys Ser Thr Pro Leu Ser  
 580 585 590  
 Gln Asn Tyr Ala Leu Thr Pro Leu Ala Ser Asp Leu Glu Asp Leu Ala  
 595 600 605  
 Leu Glu Pro Trp Ser Thr Pro Asn Thr Pro Val Ala Gly Thr Ala Glu  
 610 615 620  
 Thr Gln Asn Thr Gly Glu Ala Gly Ser Lys Ala Cys Gln Asp Gly Gln  
 625 630 635 640  
 Leu Ser Pro Thr Trp Ser Glu Ile Glu Glu Asp Leu Arg Ala Cys Phe  
 645 650 655  
 Gly Ala Glu Pro Leu Lys Lys Asp Phe Ser Glu Pro Leu Asn Leu Asp  
 660 665 670

&lt;210&gt; 11

&lt;211&gt; 60

&lt;212&gt; DNA

&lt;213&gt; part of Parvovirus NS1 variant

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (1)..(60)

&lt;400&gt; 11

att tgc tgt gtt tta aac aga caa gga ggc aaa aga aat gct gtt tta  
 Ile Cys Cys Val Leu Asn Arg Gln Gly Gly Lys Arg Asn Ala Val Leu  
 1 5 10 15

48

ttt cat gga cca  
Phe His Gly Pro  
20

<210> 12

<211> 2019

<212> DNA

<213> Parvovirus NS1 variant

<220>

<221> CDS

<222> (1)..(2016)

<400> 12

atg gct gga aat gct tac tct gat gaa gtt ttg gga gca acc aac tgg	48
Met Ala Gly Asn Ala Tyr Ser Asp Glu Val Leu Gly Ala Thr Asn Trp	
1 5 10 15	
tta aag gaa aaa agt aac cag gaa gtg ttc tca ttt gtt ttt aaa aat	96
Leu Lys Glu Lys Ser Asn Gln Glu Val Phe Ser Phe Val Phe Lys Asn	
20 25 30	
gaa aat gtt caa ctg aat gga aaa gat atc gga tgg aat agt tac aaa	144
Glu Asn Val Gln Leu Asn Gly Lys Asp Ile Gly Trp Asn Ser Tyr Lys	
35 40 45	
aaa gag ctg cag gag gac gag ctg aaa tct tta caa cga gga gcg gaa	192
Lys Glu Leu Gln Glu Asp Glu Leu Lys Ser Leu Gln Arg Gly Ala Glu	
50 55 60	
act act tgg gac caa agc gag gac atg gaa tgg gaa acc aca gtg gat	240
Thr Thr Trp Asp Gln Ser Glu Asp Met Glu Trp Glu Thr Thr Val Asp	
65 70 75 80	
gaa atg acc aaa aag caa gta ttc att ttt gat tct ttg gtt aaa aaa	288
Glu Met Thr Lys Lys Gln Val Phe Ile Phe Asp Ser Leu Val Lys Lys	
85 90 95	
tgt tta ttt gaa gtg ctt aac aca aag aat ata ttt cct ggt gat gtt	336
Cys Leu Phe Glu Val Leu Asn Thr Lys Asn Ile Phe Pro Gly Asp Val	
100 105 110	
aat tgg ttt gtg caa cat gaa tgg gga aaa gac caa ggc tgg cac tgc	384
Asn Trp Phe Val Gln His Glu Trp Gly Lys Asp Gln Gly Trp His Cys	
115 120 125	
cat gta cta att gga gga aag gac ttt agt caa gct caa ggg aaa tgg	432
His Val Leu Ile Gly Gly Lys Asp Phe Ser Gln Ala Gln Gly Lys Trp	
130 135 140	
tgg aga agg caa cta aat gtt tac tgg agc aga tgg ttg gta aca gcc	480
Trp Arg Arg Gln Leu Asn Val Tyr Trp Ser Arg Trp Leu Val Thr Ala	
145 150 155 160	
tgt aat gtg caa cta aca cca gct gaa aga att aaa cta aga gaa ata	528
Cys Asn Val Gln Leu Thr Pro Ala Glu Arg Ile Lys Leu Arg Glu Ile	
165 170 175	
gca gaa gac aat gag tgg gtt act cta ctt act tat aag cat aag caa	576
Ala Glu Asp Asn Glu Trp Val Thr Leu Leu Thr Tyr Lys His Lys Gln	
180 185 190	

acc aaa aaa gac tat acc aag tgt gtt ctt ttt gga aac atg att gct Thr Lys Lys Asp Tyr Thr Lys Cys Val Leu Phe Gly Asn Met Ile Ala 195 200 205	624
tac tat ttt tta act aaa aag aaa ata agc act agt cca cca aga gac Tyr Tyr Phe Leu Thr Lys Lys Ile Ser Thr Ser Pro Pro Arg Asp 210 215 220	672
gga ggc tat ttt ctt agc agt gac tct ggc tgg aaa act aac ttt tta Gly Gly Tyr Phe Leu Ser Ser Asp Ser Gly Trp Lys Thr Asn Phe Leu 225 230 235 240	720
aaa gaa ggc gag cgc cat cta gtg agc aaa cta tac act gat gac atg Lys Glu Gly Glu Arg His Leu Val Ser Lys Leu Tyr Thr Asp Asp Met 245 250 255	768
cgg cca gaa acg gtt gaa acc aca gta acc act gcg cag gaa act aag Arg Pro Glu Thr Val Glu Thr Thr Val Thr Thr Ala Gln Glu Thr Lys 260 265 270	816
cgc ggc aga att caa act aaa aaa gaa gtt tct att aaa act aca ctt Arg Gly Arg Ile Gln Thr Lys Lys Glu Val Ser Ile Lys Thr Thr Leu 275 280 285	864
aaa gag ctg gtg cat aaa aga gta acc tca cca gag gac tgg atg atg Lys Glu Leu Val His Lys Arg Val Thr Ser Pro Glu Asp Trp Met Met 290 295 300	912
atg cag cca gac agt tac att gaa atg atg gct caa cca ggt gga gaa Met Gln Pro Asp Ser Tyr Ile Glu Met Met Ala Gln Pro Gly Gly Glu 305 310 315 320	960
aac ctg ctg aaa aat acg cta gag att tgt aca cta act cta gcc aga Asn Leu Leu Lys Asn Thr Leu Glu Ile Cys Thr Leu Thr Leu Ala Arg 325 330 335	1008
acc aaa aca gca ttt gac tta att tta gaa aaa gct gaa acc agc aaa Thr Lys Thr Ala Phe Asp Leu Ile Leu Glu Lys Ala Glu Thr Ser Lys 340 345 350	1056
cta acc aac ttt tca ctg cct gac aca aga acc tgc aga att ttt gct Leu Thr Asn Phe Ser Leu Pro Asp Thr Arg Thr Cys Arg Ile Phe Ala 355 360 365	1104
ttt cat ggc tgg aac tat gtt aaa gtt tgc cat gct att tgc tgt gtt Phe His Gly Trp Asn Tyr Val Lys Val Cys His Ala Ile Cys Cys Val 370 375 380	1152
tta aac aga caa gga ggc aaa aga aat gct gtt tta ttt cat gga cca Leu Asn Arg Gln Gly Lys Arg Asn Ala Val Leu Phe His Gly Pro 385 390 395 400	1200
gcc agc aca ggc aaa tct att att gca caa gcc ata gca caa gca gtt Ala Ser Thr Gly Lys Ser Ile Ile Ala Gln Ala Ile Ala Gln Ala Val 405 410 415	1248
ggc aat gtt ggt tgc tat aat gca gcc aat gta aac ttt cca ttt aat Gly Asn Val Gly Cys Tyr Asn Ala Ala Asn Val Asn Phe Pro Phe Asn 420 425 430	1296
gac tgt acc aac aag aac ttg att tgg gta gaa gaa gct ggt aac ttt Asp Cys Thr Asn Lys Asn Leu Ile Trp Val Glu Glu Ala Gly Asn Phe 435 440 445	1344

gga cag caa gta aac cag ttt aaa gcc att tgc tct ggt caa act att	1392
Gly Gln Gln Val Asn Gln Phe Lys Ala Ile Cys Ser Gly Gln Thr Ile	
450 455 460	
cgc att gat caa aaa gga aaa ggc agc aaa cag att gaa cca aca cca	1440
Arg Ile Asp Gln Lys Gly Lys Gly Ser Lys Gln Ile Glu Pro Thr Pro	
465 470 475 480	
gtc atc atg acc aca aat gag aac att aca gtg gtc aga ata ggc tgc	1488
Val Ile Met Thr Thr Asn Glu Asn Ile Thr Val Val Arg Ile Gly Cys	
485 490 495	
gaa gaa aga cca gaa cac act caa cca atc aga gac aga atg ctt aac	1536
Glu Glu Arg Pro Glu His Thr Gln Pro Ile Arg Asp Arg Met Leu Asn	
500 505 510	
att cat cta aca cat acc ttg cct ggt gac ttt ggt ttg gtt gac aaa	1584
Ile His Leu Thr His Thr Leu Pro Gly Asp Phe Gly Leu Val Asp Lys	
515 520 525	
aat gaa tgg ccc atg att tgt gct tgg ttg gta aag aat ggt tac caa	1632
Asn Glu Trp Pro Met Ile Cys Ala Trp Leu Val Lys Asn Gly Tyr Gln	
530 535 540	
tct acc atg gca agc tac tgt gct aaa tgg ggc aaa gtt cct gat tgg	1680
Ser Thr Met Ala Ser Tyr Cys Ala Lys Trp Gly Lys Val Pro Asp Trp	
545 550 555 560	
tca gaa aac tgg gcg gag cca aag gtg cca act cct ata aat tta cta	1728
Ser Glu Asn Trp Ala Glu Pro Lys Val Pro Thr Pro Ile Asn Leu Leu	
565 570 575	
ggt tcg gca cgc tca cca ttc acg aca ccg aaa agt acg cct ctc agc	1776
Gly Ser Ala Arg Ser Pro Phe Thr Thr Pro Lys Ser Thr Pro Leu Ser	
580 585 590	
cag aac tat gca cta act cca ctt gca tcg gat ctc gag gac ctg gct	1824
Gln Asn Tyr Ala Leu Thr Pro Leu Ala Ser Asp Leu Glu Asp Leu Ala	
595 600 605	
tta gag cct tgg agc aca cca aat act cct gtt gcg ggc act gca gaa	1872
Leu Glu Pro Trp Ser Thr Pro Asn Thr Pro Val Ala Gly Thr Ala Glu	
610 615 620	
acc cag aac act ggg gaa gct ggt tcc aaa gcc tgc caa gat ggt caa	1920
Thr Gln Asn Thr Gly Glu Ala Gly Ser Lys Ala Cys Gln Asp Gly Gln	
625 630 635 640	
ctg agc cca act tgg tca gag atc gag gag gat ttg aga gcg tgc ttc	1968
Leu Ser Pro Thr Trp Ser Glu Ile Glu Glu Asp Leu Arg Ala Cys Phe	
645 650 655	
ggt gcg gaa ccg ttg aag aaa gac ttc agc gag ccg ctg aac ttg gac	2016
Gly Ala Glu Pro Leu Lys Lys Asp Phe Ser Glu Pro Leu Asn Leu Asp	
660 665 670	
taa	2019

&lt;210&gt; 13

&lt;211&gt; 20

&lt;212&gt; PRT

&lt;213&gt; part of Parvovirus NS1 variant

&lt;400&gt; 13



20

Ile Cys Cys Val Leu Asn Arg Gln Gly Gly Lys Arg Asn Ala Val Leu  
 1 5 10 15

Phe His Gly Pro  
 20

<210> 14  
 <211> 672  
 <212> PRT  
 <213> Parvovirus NS1 variant

<400> 14

Met Ala Gly Asn Ala Tyr Ser Asp Glu Val Leu Gly Ala Thr Asn Trp  
 1 5 10 15

Leu Lys Glu Lys Ser Asn Gln Glu Val Phe Ser Phe Val Phe Lys Asn  
 20 25 30

Glu Asn Val Gln Leu Asn Gly Lys Asp Ile Gly Trp Asn Ser Tyr Lys  
 35 40 45

Lys Glu Leu Gln Glu Asp Glu Leu Lys Ser Leu Gln Arg Gly Ala Glu  
 50 55 60

Thr Thr Trp Asp Gln Ser Glu Asp Met Glu Trp Glu Thr Thr Val Asp  
 65 70 75 80

Glu Met Thr Lys Lys Gln Val Phe Ile Phe Asp Ser Leu Val Lys Lys  
 85 90 95

Cys Leu Phe Glu Val Leu Asn Thr Lys Asn Ile Phe Pro Gly Asp Val  
 100 105 110

Asn Trp Phe Val Gln His Glu Trp Gly Lys Asp Gln Gly Trp His Cys  
 115 120 125

His Val Leu Ile Gly Gly Lys Asp Phe Ser Gln Ala Gln Gly Lys Trp  
 130 135 140

Trp Arg Arg Gln Leu Asn Val Tyr Trp Ser Arg Trp Leu Val Thr Ala  
 145 150 155 160

Cys Asn Val Gln Leu Thr Pro Ala Glu Arg Ile Lys Leu Arg Glu Ile  
 165 170 175

Ala Glu Asp Asn Glu Trp Val Thr Leu Leu Thr Tyr Lys His Lys Gln  
 180 185 190

Thr Lys Lys Asp Tyr Thr Lys Cys Val Leu Phe Gly Asn Met Ile Ala  
 195 200 205

Tyr Tyr Phe Leu Thr Lys Lys Lys Ile Ser Thr Ser Pro Pro Arg Asp  
 210 215 220

Gly Gly Tyr Phe Leu Ser Ser Asp Ser Gly Trp Lys Thr Asn Phe Leu  
 225 230 235 240

Lys Glu Gly Glu Arg His Leu Val Ser Lys Leu Tyr Thr Asp Asp Met  
 245 250 255

Arg Pro Glu Thr Val Glu Thr Thr Val Thr Thr Ala Gln Glu Thr Lys  
 260 265 270

Arg Gly Arg Ile Gln Thr Lys Lys Glu Val Ser Ile Lys Thr Thr Leu  
 275 280 285  
 Lys Glu Leu Val His Lys Arg Val Thr Ser Pro Glu Asp Trp Met Met  
 290 300  
~~Met Gln Pro Asp Ser Tyr Ile Glu Met Met Ala Gln Pro Gly Gly Glu~~  
~~305 310 315 320~~  
 Asn Leu Leu Lys Asn Thr Leu Glu Ile Cys Thr Leu Thr Leu Ala Arg  
 325 330 335  
 Thr Lys Thr Ala Phe Asp Leu Ile Leu Glu Lys Ala Glu Thr Ser Lys  
 340 345 350  
 Leu Thr Asn Phe Ser Leu Pro Asp Thr Arg Thr Cys Arg Ile Phe Ala  
 355 360 365  
 Phe His Gly Trp Asn Tyr Val Lys Val Cys His Ala Ile Cys Cys Val  
 370 375 380  
 Leu Asn Arg Gln Gly Gly Lys Arg Asn Ala Val Leu Phe His Gly Pro  
 385 390 395 400  
 Ala Ser Thr Gly Lys Ser Ile Ile Ala Gln Ala Ile Ala Gln Ala Val  
 405 410 415  
 Gly Asn Val Gly Cys Tyr Asn Ala Ala Asn Val Asn Phe Pro Phe Asn  
 420 425 430  
 Asp Cys Thr Asn Lys Asn Leu Ile Trp Val Glu Glu Ala Gly Asn Phe  
 435 440 445  
 Gly Gln Gln Val Asn Gln Phe Lys Ala Ile Cys Ser Gly Gln Thr Ile  
 450 455 460  
 Arg Ile Asp Gln Lys Gly Lys Gly Ser Lys Gln Ile Glu Pro Thr Pro  
 465 470 475 480  
 Val Ile Met Thr Thr Asn Glu Asn Ile Thr Val Val Arg Ile Gly Cys  
 485 490 495  
 Glu Glu Arg Pro Glu His Thr Gln Pro Ile Arg Asp Arg Met Leu Asn  
 500 505 510  
 Ile His Leu Thr His Thr Leu Pro Gly Asp Phe Gly Leu Val Asp Lys  
 515 520 525  
 Asn Glu Trp Pro Met Ile Cys Ala Trp Leu Val Lys Asn Gly Tyr Gln  
 530 535 540  
 Ser Thr Met Ala Ser Tyr Cys Ala Lys Trp Gly Lys Val Pro Asp Trp  
 545 550 555 560  
 Ser Glu Asn Trp Ala Glu Pro Lys Val Pro Thr Pro Ile Asn Leu Leu  
 565 570 575  
 Gly Ser Ala Arg Ser Pro Phe Thr Thr Pro Lys Ser Thr Pro Leu Ser  
 580 585 590  
 Gln Asn Tyr Ala Leu Thr Pro Leu Ala Ser Asp Leu Glu Asp Leu Ala  
 595 600 605  
 Leu Glu Pro Trp Ser Thr Pro Asn Thr Pro Val Ala Gly Thr Ala Glu  
 610 615 620

22

Thr Gln Asn Thr Gly Glu Ala Gly Ser Lys Ala Cys Gln Asp Gly Gln  
625 630 635 640

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Gly Ala Glu Pro Leu Lys Lys Asp Phe Ser Glu Pro Leu Asn Leu Asp  
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tta aag gaa aaa agt aac cag gaa gtg ttc tca ttt gtt ttt aaa aat 96  
Leu Lys Glu Lys Ser Asn Gln Glu Val Phe Ser Phe Val Phe Lys Asn  
20 25 30

gaa aat gtt caa ctg aat gga aaa gat atc gga tgg aat agt tac aaa 144  
Glu Asn Val Gln Leu Asn Gly Lys Asp Ile Gly Trp Asn Ser Tyr Lys  
35 40 45

aaa gag ctg cag gag gac gag ctg aaa tct tta caa cga gga gcg gaa 192  
Lys Glu Leu Gln Glu Asp Glu Leu Lys Ser Leu Gln Arg Gly Ala Glu  
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act act tgg gac caa agc gag gac atg gaa tgg gaa acc aca gtg gat 240  
Thr Thr Trp Asp Gln Ser Glu Asp Met Glu Trp Glu Thr Thr Val Asp  
65 70 75 80

gaa atg acc aaa aag caa gta ttc att ttt gat tct ttg gtt aaa aaa	288
Glu Met Thr Lys Lys Gln Val Phe Ile Phe Asp Ser Leu Val Lys Lys	
85 90 95	
tgt tta ttt gaa gtg ctt aac aca aag aat ata ttt cct ggt gat gtt	336
Cys Leu Phe Glu Val Leu Asn Thr Lys Asn Ile Phe Pro Gly Asp Val	
100 105 110	
aat tgg ttt gtg caa cat gaa tgg gga aaa gac caa ggc tgg cac tgc	384
Asn Trp Phe Val Gln His Glu Trp Gly Lys Asp Gln Gly Trp His Cys	
115 120 125	
cat gta cta att gga gga aag gac ttt agt caa gct caa ggg aaa tgg	432
His Val Leu Ile Gly Gly Lys Asp Phe Ser Gln Ala Gln Gly Lys Trp	
130 135 140	
tgg aga agg caa cta aat gtt tac tgg agc aga tgg ttg gta aca gcc	480
Trp Arg Arg Gln Leu Asn Val Tyr Trp Ser Arg Trp Leu Val Thr Ala	
145 150 155 160	
tgt aat gtg caa cta aca cca gct gaa aga att aaa cta aga gaa ata	528
Cys Asn Val Gln Leu Thr Pro Ala Glu Arg Ile Lys Leu Arg Glu Ile	
165 170 175	
gca gaa gac aat gag tgg gtt act cta ctt act tat aag cat aag caa	576
Ala Glu Asp Asn Glu Trp Val Thr Leu Leu Thr Tyr Lys His Lys Gln	
180 185 190	
acc aaa aaa gac tat acc aag tgt gtt ctt ttt gga aac atg att gct	624
Thr Lys Lys Asp Tyr Thr Lys Cys Val Leu Phe Gly Asn Met Ile Ala	
195 200 205	
tac tat ttt tta act aaa aag aaa ata agc act agt cca cca aga gac	672
Tyr Tyr Phe Leu Thr Lys Lys Lys Ile Ser Thr Ser Pro Pro Arg Asp	
210 215 220	
gga ggc tat ttt ctt agc agt gac tct ggc tgg aaa act aac ttt tta	720
Gly Gly Tyr Phe Leu Ser Ser Asp Ser Gly Trp Lys Thr Asn Phe Leu	
225 230 235 240	
aaa gaa ggc gag cgc cat cta gtg agc aaa cta tac act gat gac atg	768
Lys Glu Gly Glu Arg His Leu Val Ser Lys Leu Tyr Thr Asp Asp Met	
245 250 255	
cgg cca gaa acg gtt gaa acc aca gta acc act gcg cag gaa act aag	816
Arg Pro Glu Thr Val Glu Thr Thr Val Thr Thr Ala Gln Glu Thr Lys	
260 265 270	
cgc ggc aga att caa act aaa aaa gaa gtt tct att aaa act aca ctt	864
Arg Gly Arg Ile Gln Thr Lys Lys Glu Val Ser Ile Lys Thr Thr Leu	
275 280 285	
aaa gag ctg gtg cat aaa aga gta acc tca cca gag gac tgg atg atg	912
Lys Glu Leu Val His Lys Arg Val Thr Ser Pro Glu Asp Trp Met Met	
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Met Gln Pro Asp Ser Ile Glu Met Met Ala Gln Pro Gly Gly Glu	
305 310 315 320	
aac ctg ctg aaa aat acg cta gag att tgt aca cta act cta gcc aga	1008
Asn Leu Leu Lys Asn Thr Leu Glu Ile Cys Thr Leu Thr Leu Ala Arg	
325 330 335	
acc aaa aca gca ttt gac tta att tta gaa aaa gct gaa acc agc aaa	1056
Thr Lys Thr Ala Phe Asp Leu Ile Leu Glu Lys Ala Glu Thr Ser Lys	
340 345 350	

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gcc agc aca ggc aaa tct att att gca caa gcc ata gca caa gca gtt Ala Ser Thr Gly Lys Ser Ile Ile Ala Gln Ala Ile Ala Gln Ala Val 405 410 415	1248
ggc aat gtt ggt tgc tat aat gca gcc aat gta aac ttt cca ttt aat Gly Asn Val Gly Cys Tyr Asn Ala Ala Asn Val Asn Phe Pro Phe Asn 420 425 430	1296
gac tgt acc aac aag aac ttg att tgg gta gaa gaa gct ggt aac ttt Asp Cys Thr Asn Lys Asn Leu Ile Trp Val Glu Glu Ala Gly Asn Phe 435 440 445	1344
gga cag caa gta aac cag ttt aaa gcc att tgc tct ggt caa gct att Gly Gln Gln Val Asn Gln Phe Lys Ala Ile Cys Ser Gly Gln Ala Ile 450 455 460	1392
cgc att gat caa aaa gga aaa ggc agc aaa cag att gaa cca aca cca Arg Ile Asp Gln Lys Gly Lys Gly Ser Lys Gln Ile Glu Pro Thr Pro 465 470 475 480	1440
gtc atc atg acc aca aat gag aac att aca gtg gtc aga ata ggc tgc Val Ile Met Thr Asn Glu Asn Ile Thr Val Val Arg Ile Gly Cys 485 490 495	1488
gaa gaa aga cca gaa cac act caa cca atc aga gac aga atg ctt aac Glu Glu Arg Pro Glu His Thr Gln Pro Ile Arg Asp Arg Met Leu Asn 500 505 510	1536
att cat cta aca cat acc ttg cct ggt gac ttt ggt ttg gtt gac aaa Ile His Leu Thr His Thr Leu Pro Gly Asp Phe Gly Leu Val Asp Lys 515 520 525	1584
aat gaa tgg ccc atg att tgt gct tgg ttg gta aag aat ggt tac caa Asn Glu Trp Pro Met Ile Cys Ala Trp Leu Val Lys Asn Gly Tyr Gln 530 535 540	1632
tct acc atg gca agc tac tgt gct aaa tgg ggc aaa gtt cct gat tgg Ser Thr Met Ala Ser Tyr Cys Ala Lys Trp Gly Lys Val Pro Asp Trp 545 550 555 560	1680
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ggt tcg gca cgc tca cca ttc acg aca ccg aaa agt acg cct ctc agc Gly Ser Ala Arg Ser Pro Phe Thr Thr Pro Lys Ser Thr Pro Leu Ser 580 585 590	1776
cag aac tat gca cta act cca ctt gca tcg gat ctc gag gac ctg gct Gln Asn Tyr Ala Leu Thr Pro Leu Ala Ser Asp Leu Glu Asp Leu Ala 595 600 605	1824
tta gag cct tgg agc aca cca aat act cct gtt gcg ggc act gca gaa Leu Glu Pro Trp Ser Thr Pro Asn Thr Pro Val Ala Gly Thr Ala Glu 610 615 620	1872

25

acc cag aac act ggg gaa gct ggt tcc aaa gcc tgc caa gat ggt caa 1920  
 Thr Gln Asn Thr Gly Glu Ala Gly Ser Lys Ala Cys Gln Asp Gly Gln  
 625 630 635 640

ctg agc cca act tgg tca gag atc gag gag gat ttg aga gcg tgc ttc 1968  
 Leu Ser Pro Thr Trp Ser Glu Ile Glu Glu Asp Leu Arg Ala Cys Phe  
 645 650 655

ggt gcg gaa ccg ttg aag aaa gac ttc agc gag ccg ctg aac ttg gac 2016  
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taa 2019

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 <212> PRT  
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<210> 18  
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Glu Asn Val Gln Leu Asn Gly Lys Asp Ile Gly Trp Asn Ser Tyr Lys  
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Lys Glu Leu Gln Glu Asp Glu Leu Lys Ser Leu Gln Arg Gly Ala Glu  
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Thr Thr Trp Asp Gln Ser Glu Asp Met Glu Trp Glu Thr Thr Val Asp  
 65 70 75 80

Glu Met Thr Lys Lys Gln Val Phe Ile Phe Asp Ser Leu Val Lys Lys  
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Cys Leu Phe Glu Val Leu Asn Thr Lys Asn Ile Phe Pro Gly Asp Val  
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Asn Trp Phe Val Gln His Glu Trp Gly Lys Asp Gln Gly Trp His Cys  
 115 120 125

His Val Leu Ile Gly Gly Lys Asp Phe Ser Gln Ala Gln Gly Lys Trp  
 130 135 140

Trp 145	Arg	Arg	Gln	Leu	Asn 150	Val	Tyr	Trp	Ser	Arg 155	Trp	Leu	Val	Thr	Ala 160
Cys	Asn	Val	Gln	Leu 165	Thr	Pro	Ala	Glu	Arg 170	Ile	Lys	Leu	Arg	Glu 175	Ile
Ala	Glu	Asp	Asn 180	Glu	Trp	Val	Thr	Leu 185	Leu	Thr	Tyr	Lys	His 190	Lys	Gln
Thr	Lys 195	Lys	Asp	Tyr	Thr	Lys	Cys 200	Val	Leu	Phe	Gly	Asn 205	Met	Ile	Ala
Tyr 210	Tyr	Phe	Leu	Thr	Lys 215	Lys	Lys	Ile	Ser	Thr	Ser 220	Pro	Pro	Arg	Asp
Gly 225	Gly	Tyr	Phe	Leu	Ser 230	Ser	Asp	Ser	Gly	Trp 235	Lys	Thr	Asn	Phe	Leu 240
Lys	Glu	Gly	Glu	Arg 245	His	Leu	Val	Ser	Lys 250	Leu	Tyr	Thr	Asp	Asp 255	Met
Arg	Pro	Glu	Thr 260	Val	Glu	Thr	Thr	Val 265	Thr	Thr	Ala	Gln	Glu 270	Thr	Lys
Arg	Gly	Arg 275	Ile	Gln	Thr	Lys	Lys 280	Glu	Val	Ser	Ile	Lys 285	Thr	Thr	Leu
Lys 290	Glu	Leu	Val	His	Lys	Arg 295	Val	Thr	Ser	Pro	Glu 300	Asp	Trp	Met	Met
Met 305	Gln	Pro	Asp	Ser	Tyr 310	Ile	Glu	Met	Met	Ala 315	Gln	Pro	Gly	Gly	Glu 320
Asn	Leu	Leu	Lys	Asn 325	Thr	Leu	Glu	Ile	Cys 330	Thr	Leu	Thr	Leu	Ala 335	Arg
Thr	Lys	Thr	Ala 340	Phe	Asp	Leu	Ile	Leu 345	Glu	Lys	Ala	Glu	Thr 350	Ser	Lys
Leu	Thr	Asn 355	Phe	Ser	Leu	Pro	Asp 360	Thr	Arg	Thr	Cys	Arg 365	Ile	Phe	Ala
Phe 370	His	Gly	Trp	Asn	Tyr	Val 375	Lys	Val	Cys	His	Ala 380	Ile	Cys	Cys	Val
Leu 385	Asn	Arg	Gln	Gly	Gly 390	Lys	Arg	Asn	Thr	Val 395	Leu	Phe	His	Gly	Pro 400
Ala	Ser	Thr	Gly	Lys 405	Ser	Ile	Ile	Ala	Gln 410	Ala	Ile	Ala	Gln	Ala 415	Val
Gly	Asn	Val	Gly 420	Cys	Tyr	Asn	Ala	Ala 425	Asn	Val	Asn	Phe	Pro 430	Phe	Asn
Asp	Cys	Thr 435	Asn	Lys	Asn	Leu	Ile 440	Trp	Val	Glu	Glu	Ala 445	Gly	Asn	Phe
Gly 450	Gln	Gln	Val	Asn	Gln	Phe 455	Lys	Ala	Ile	Cys	Ser 460	Gly	Gln	Ala	Ile
Arg 465	Ile	Asp	Gln	Lys	Gly 470	Lys	Gly	Ser	Lys	Gln 475	Ile	Glu	Pro	Thr	Pro 480
Val	Ile	Met	Thr	Thr 485	Asn	Glu	Asn	Ile	Thr 490	Val	Val	Arg	Ile	Gly 495	Cys
Glu	Glu	Arg	Pro	Glu	His	Thr	Gln	Pro	Ile	Arg	Asp	Arg	Met	Leu	Asn

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Asn	Glu	Trp	Pro	Met	Ile	Cys	Ala	Trp	Leu	Val	Lys	Asn	Gly	Tyr	Gln
	530					535					540				
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Ser	Glu	Asn	Trp	Ala	Glu	Pro	Lys	Val	Pro	Thr	Pro	Ile	Asn	Leu	Leu
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Gly	Ser	Ala	Arg	Ser	Pro	Phe	Thr	Thr	Pro	Lys	Ser	Thr	Pro	Leu	Ser
			580					585					590		
Gln	Asn	Tyr	Ala	Leu	Thr	Pro	Leu	Ala	Ser	Asp	Leu	Glu	Asp	Leu	Ala
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Gly	Ala	Glu	Pro	Leu	Lys	Lys	Asp	Phe	Ser	Glu	Pro	Leu	Asn	Leu	Asp
			660					665					670		



# INTERNATIONAL SEARCH REPORT

International Application No

PCT/EP 00/07835

## A. CLASSIFICATION OF SUBJECT MATTER

IPC 7 C12N15/35 C07K14/015 C07K16/08 G01N33/569 C12Q1/70  
A61K35/76 A61K48/00

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 7 C12N C07K A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

EPO-Internal, STRAND, MEDLINE, BIOSIS

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	D LEGENDRE & J ROMMELAERE: "Terminal regions of the NS-1 protein of the parvovirus Minute Virus of Mice are involved in cytotoxicity and promoter trans inhibition" JOURNAL OF VIROLOGY, vol. 66, no. 10, October 1992 (1992-10), pages 5705-5713, XP000867510 AMERICAN SOCIETY FOR MICROBIOLOGY US *mutants pMMBa131 and pULB3201; figure 1 and page 5710 first paragraph*  --- -/-	1-5,7-11

☒ Further documents are listed in the continuation of box C.

☐ Patent family members are listed in annex.

### \* Special categories of cited documents :

- \*A\* document defining the general state of the art which is not considered to be of particular relevance
- \*E\* earlier document but published on or after the international filing date
- \*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- \*O\* document referring to an oral disclosure, use, exhibition or other means
- \*P\* document published prior to the international filing date but later than the priority date claimed

- \*T\* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
- \*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
- \*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.
- \*G\* document member of the same patent family

Date of the actual completion of the international search

19 January 2001

Date of mailing of the international search report

30.01.01

Name and mailing address of the ISA

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Authorized officer

Cupido, M

## INTERNATIONAL SEARCH REPORT

International Application No

PCT/EP 00/07835

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	LI X ET AL: "Mutation of lysine 405 to serine in the parvovirus H-1 NS1 abolishes its function for viral DNA replication, late promoter activation, and cytotoxicity" JOURNAL OF VIROLOGY, vol. 64, no. 10, October 1990 (1990-10), pages 4654-4660, XP000867496 AMERICAN SOCIETY FOR MICROBIOLOGY US page 4656 -page 4657	1,3,7, 9-11
A	J P F NÜESCH ET AL: "Sequence motifs in the replicator protein of parvovirus MVM essential for nicking and covalent attachment to the viral origin: identification of the linking tyrosine" VIROLOGY,US,ACADEMIC PRESS,ORLANDO, vol. 209, no. 1, 10 May 1995 (1995-05-10), pages 122-135, XP002088311 ISSN: 0042-6822 page 127 -page 131	1-13
A	S F COTTMORE ET AL: "The NS1 polypeptide of the murine parvovirus MVM binds to DNA sequences containing the motif (ACCA)2-3" JOURNAL OF VIROLOGY,US,THE AMERICAN SOCIETY FOR MICROBIOLOGY, vol. 69, no. 3, pages 1652-1660, XP002088309 ISSN: 0022-538X page 1658, left-hand column, last paragraph -right-hand column	1-13

# INTERNATIONAL SEARCH REPORT

International application No.  
PCT/EP 00/07835

## Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:  
~~Although claims 14 and 15 are directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the NS1 variant.~~
2. ☐ Claims Nos.:  
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

### Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.